

(19) World Intellectual Property Organization
International Bureau(43) International Publication Date
21 November 2002 (21.11.2002)

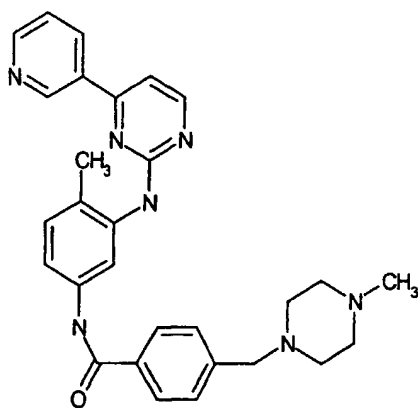
PCT

(10) International Publication Number
WO 02/093164 A2(51) International Patent Classification⁷: **G01N 33/48**(74) Agents: **LEIDESCHER, Thomas** et al.; Zimmermann & Partner, Postfach 330 920, 80069 München (DE).(21) International Application Number: **PCT/EP02/05420**(22) International Filing Date: **16 May 2002 (16.05.2002)**(25) Filing Language: **English**(26) Publication Language: **English**(30) Priority Data:
01111858.5 16 May 2001 (16.05.2001) EP
60/293,528 29 May 2001 (29.05.2001) US
01117113.9 13 July 2001 (13.07.2001) EP
60/305,898 18 July 2001 (18.07.2001) US(81) Designated States (*national*): AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, BZ, CA, CH, CN, CO, CR, CU, CZ, DE, DK, DM, DZ, EC, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NO, NZ, OM, PH, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, US, UZ, VN, YU, ZA, ZM, ZW.(84) Designated States (*regional*): ARIPO patent (GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZM, ZW), Eurasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European patent (AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, TR), OAPI patent (BF, BJ, CF, CG, CI, CM, GA, GN, GQ, GW, ML, MR, NE, SN, TD, TG).(71) Applicant (*for all designated States except US*): **AXXIMA PHARMACEUTICALS AG** [DE/DE]; Am Klopferspitz 19, 82152 Martinsried (DE).

(72) Inventors; and

(75) Inventors/Applicants (*for US only*): **STEIN-GERLACH, Matthias** [DE/DE]; Stockdorfer Strasse 38A, 81475 München (DE). **SALASSIDIS, Konstadinos** [GR/DE]; Echinger strasse 20, 85386 Echting (DE). **BACHER, Gerald** [DE/DE]; Kriegerstrasse 62, 82110 Germering (DE). **MÜLLER, Stefan** [DE/DE]; Thalkirchner Str. 184, 81371 München (DE).

Published:

— *without international search report and to be republished upon receipt of that report**For two-letter codes and other abbreviations, refer to the "Guidance Notes on Codes and Abbreviations" appearing at the beginning of each regular issue of the PCT Gazette.*(54) Title: **PYRIDYLPYRIMIDINE DERIVATIVES AS EFFECTIVE COMPOUNDS AGAINST PRION DISEASES**

Compound 53 (Gleevec™)

(57) Abstract: The present invention relates to pyridylpyrimidine derivatives of the general formula (I): wherein R represents hydrogen or methyl and Z represents nitrogen containing functional groups, the use of the pyridylpyrimidine derivatives as pharmaceutically active agents, especially for the prophylaxis and/or treatment of prion infections and prion diseases, as well as compositions containing at least one pyridylpyrimidine derivative and/or pharmaceutically acceptable salt thereof. Furthermore, the present invention is directed to methods for preventing and/or treating prion infections and prion diseases using said pyridylpyrimidine derivatives. Human cellular protein kinases, phosphatases and cellular signal transduction molecules are disclosed as targets for detecting, preventing and/or treating prion infections and diseases, especially BSE, vCJD, or CJD, which can be inhibited by the inventive pyridylpyrimidine derivatives.

**Pyridylpyrimidine derivatives as effective compounds against
prion infections and prion diseases**

5

Specification

The present invention relates to pyridylpyrimidine derivatives, the use of the pyridylpyrimidine derivatives as pharmaceutically active agents, especially for the prophylaxis and/or treatment of prion infections and prion diseases, as well as compositions containing at least one pyridylpyrimidine derivative and/or pharmaceutically acceptable salt thereof, and methods for preventing and/or treating prion infections and prion diseases. Furthermore, human cellular protein kinases, phosphatases and cellular signal transduction molecules are disclosed as targets for detecting, preventing and/or treating prion infections and diseases, especially BSE, vCJD, or CJD.

Background of the invention

Pyridylpyrimidine derivatives are known from WO 9509851 as effective compounds for chemotherapy of tumors, from WO 9509853, EP-A-0 588 762, WO 9509847, WO 9903854, and EP-B-0 564 409 as effective compounds for treatment of tumors. Furthermore, EP-B-0 564 409 discloses the use of said compounds in the treatment of arteriosclerosis and Exp. Opin. Ther. Patents, 1998, 8(12), 1599-1625 describes the use of pyridylpyrimidine derivatives, especially of GleevecTM, the Novartis compound CGP 57148, as tyrosine kinase inhibitors in cancer treatment.

Prions are infectious agents which do not have a nucleic acid genome. It seems that a protein alone is the infectious agent. A prion has been defined as "small proteinaceous infectious particle which resists inactivation by procedures that modify nucleic acids". The discovery that proteins alone can transmit an infectious disease has come as a considerable surprise to the scientific community. Prion diseases are often called "transmissible spongiform encephalopathies", because of the post mortem appearance of the brain with large vacuoles in the cortex and cerebellum. Probably most mammalian species develop these diseases. Prion diseases are a group of neurodegenerative disorders of humans and animals and the prion diseases can manifest as sporadic, genetic or infectious disorders. Examples for prion diseases acquired

by exogenous infection are the Bovine spongiform encephalitis (BSE) of cattle and the new variant of Creutzfeld-Jakob disease (vCJD) caused by BSE. Further examples include kuru, Gerstmann-Sträussler-Scheinker disease of humans as well as scrapie of animals. For many years, the prion diseases were thought to be caused by viruses despite intriguing evidence to the contrary. The unique characteristic common to all of these disorders, whether sporadic, dominantly inherited, or acquired by infection, is that they involve the aberrant metabolism of the prion protein (PrP). In many cases, the cellular prion protein (PrP^c) ["c" refers to cellular] is converted into the scrapie isoform (PrP^{Sc}) ["Sc" refers to Scrapie] by a posttranslational process that involves a conformational change. Often, the human prion diseases are transmissible to experimental animals and all of the inherited prion diseases segregate with PrP gene mutations.

These prion diseases in animals and humans have a long incubation period and a long clinical course, and are always fatal leading via decerebration to death within an average period of 7 months (CJD). Neuropathological features consist of neuronal vacuolization, neuronal death and gliosis with hyperastrocytosis. The precise diagnosis of transmissible neurodegenerative diseases can be established only by the examination of the central nervous system after biopsy or autopsy.

Clinical symptoms of the disease are progressive dementia, myoclonus and prominent ataxia with the additional clinical features of dysautonomia and delirious psychomotor excitement and with relatively preserved verbal responses.

Between 1980 and, roughly, 1996, about 750,000 cattle infected with BSE were slaughtered for human consumption in Great Britain (Anderson, R. M. *et al. Nature* 382, 779-788, 1996; Ferguson, N. M., Donnelly, C. A., Woolhouse, M. E. J. & Anderson, R. M. *Phil. Trans. R. Soc. Lond. B* 352, 803-838, 1997). The annual incidence of vCJD (3, 10, 10, 18, 14 and 33 deaths in 1995–2000, respectively) can be interpreted as a first sign of a steady or exponential increase over the next years. The suggestion by the European Union Scientific Steering Committee that up to 500,000 people could have been exposed to BSE from a single infected bovine has fuelled speculation that millions of consumers are at risk.

Recent findings demonstrate that the pathogenic PrP^{Sc} of vCJD can be found in the lymph system (e.g. tonsils, lymph nodes) in humans suggesting a high risk of horizontal spread via lymph and/or blood transmission, dramatically increasing the number of people at risk.

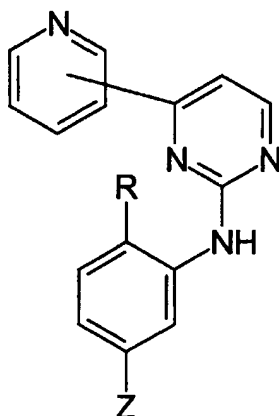
The medical need in prion diseases today can be clearly defined as the establishment of a diagnostic system, that can detect the disease as early as possible in living humans and/or animals, to estimate the medical need for the treatment in the future and to identify the infected animals to remove them from the food chain. The medical need for prion diseases in the future (approximately starting in 5-10 years) will be medical treatment that inhibits the disease symptoms, the manifestation and/or progression of the disease.

It is object of the present invention to provide novel and also known compounds which can be used as pharmaceutically active agents, especially for prophylaxis and/or treatment of prion infections and prion diseases, methods wherein said compounds are used in order to treat prion infections and prion diseases and compositions containing at least one inventive compound and/or pharmaceutically acceptable salt thereof as a pharmaceutically active ingredient.

The object of the present invention is solved by the teaching of the independent claims. Further advantageous features, aspects and details of the invention are evident from the dependent claims, the description, the examples, and the figures of the present application.

Description of the invention

One aspect of the present invention is related to compounds of the general formula (I):



wherein:

R represents hydrogen or methyl;

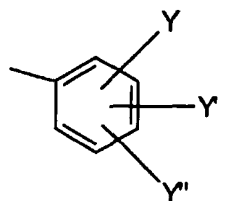
Y, Y', Y'' are independently of each other -H, -F, -Cl, -Br, -I, -CH₂F, -CH₂Cl, -CH₂Br, -CH₂I, -OH, -OCH₃, -CH₃, -CN, -OCF₃, 4-methylpiperazin-1-yl-methyl, -C(CH₃)=N-NH-C(NH)-NH₂;

Z represents -NO₂, -NH₂, -NH-CO-X, -NH-CS-X, -NH-CO-NH-X,

5 -NH-SO₂-X;

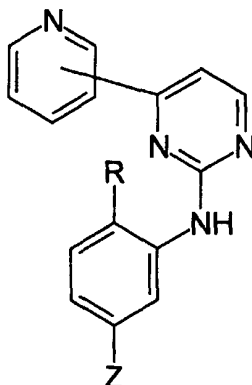
X represents thiophenyl, cyclohexyl, isoquinolinyl, naphthyl, quinolinyl,

cyclopentyl, pyridinyl, naphthyridinyl, or



and pharmaceutically acceptable salts thereof.

Another aspect of the present invention relates to the use of compounds of the
10 general formula (I):



wherein:

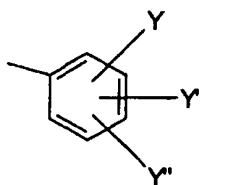
R represents hydrogen or methyl;

Y, Y', Y'' are independently of each other -H, -F, -Cl, -Br, -I, -CH₂F, -CH₂Cl, -CH₂Br, -CH₂I, -OH, -OCH₃, -CH₃, -CN, -OCF₃, 4-methylpiperazin-1-yl-methyl, -C(CH₃)=N-NH-C(NH)-NH₂;

15 Z represents -NO₂, -NH₂, -NH-CO-X, -NH-CS-X, -NH-CO-NH-X, -NH-SO₂-X;

X represents thiophenyl, cyclohexyl, isoquinolinyl, naphthyl, quinolinyl,

cyclopentyl, pyridinyl, naphthyridinyl, or



and pharmaceutically acceptable salts thereof as pharmaceutically active agents, especially for prophylaxis and/or treatment of infectious diseases, or in a more general sense, for prophylaxis and/or treatment of neurodegenerative diseases.

- 5 Thus, one embodiment of the present invention disclosed herein is directed to a method for preventing and/or treating infections and/or diseases associated with said infections in an individual. Said method comprises administering to the individual an amount of at least one compound according to general formula (I) and/or pharmaceutically acceptable salts thereof effective to prevent and/or treat
10 said infections and/or diseases. Most preferred is the administration of a compound 53.

As revealed for the first time herein, the present invention discloses the use of compounds of the general formula (I) for the prophylaxis and/or treatment of prion
15 infections and prion diseases. As described above, said pyridylpyrimidine derivatives have first of all been used in tumor therapy. The Novartis compound GleevecTM also known as GlivecTM, CGP-57148B, imatinib mesylate, STI-571, STI-571A, CAS 152459-95-5, or 4-((Methyl-1-piperazinyl)methyl)-N-[4-methyl-3-[[4-(3-pyridinyl)-2-pyrimidinyl]amino]-phenyl]benzamide methanesulfonate, has been
20 registered in many countries as anticancer drug. This GleevecTM compound (compound 53) is also the most active one in the indication prion diseases.

The name "prion" is used to describe the causative agents which underlie the transmissible spongiform encephalopathies. A prion is proposed to be a novel
25 infectious particle that differs from viruses and viroids. It is composed solely of one unique protein that resists most inactivation procedures such as heat, radiation, and proteases. The latter characteristic has led to the term protease-resistant isoform of the prion protein. The protease-resistant isoform has been proposed to slowly catalyze the conversion of the normal prion protein into the
30 abnormal form.

The term "isoform" in the context of prions means two proteins with exactly the same amino acid sequence that are folded into molecules with dramatically different tertiary structures. The normal cellular isoform of the prion protein
35 (PrP^C) has a high α -helix content, a low β -sheet content, and is sensitive to protease digestion. The abnormal, disease-causing isoform (PrP^{Sc}) has a lower α -helix content, a much higher β -sheet content, and is much more resistant to protease digestion.

Moreover, in a more general sense, the present invention is concerned with the prophylaxis and/or treatment of neurodegenerative diseases. For example, Alzheimer is a well-known neurodegenerative disease.

- 5 Preferred are the compounds wherein R represents hydrogen. Also preferred are compounds wherein Z represents $-\text{NH}-\text{CO}-\text{X}$ or $-\text{NH}-\text{SO}_2-\text{X}$ and/or wherein Y, Y', Y'' are independently of each other $-\text{H}$, $-\text{F}$, $-\text{Cl}$, $-\text{CH}_2\text{F}$, $-\text{CH}_2\text{Cl}$, $-\text{OH}$, $-\text{OCH}_3$, $-\text{CN}$, $-\text{OCF}_3$, or a 4-methylpiperazin-1-yl-methyl residue.
- 10 Also preferred are the following pyridylpyrimidine derivatives selected from the group comprising:
- Compound 1: (3-Nitrophenyl)-(4-pyridin-3-yl-pyrimidin-2-yl)-amine;
 - Compound 2: (3-Aminophenyl)-(4-pyridin-3-yl-pyrimidin-2-yl)-amine;
 - Compound 3: (5-Amino-2-methylphenyl)-(4-pyridin-3-yl-pyrimidin-2-yl)-
15 amine;
 - Compound 4: 4-Chloromethyl-N-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
 - Compound 5: 4-Chloromethyl-N-[3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
 - Compound 6: 4-(4-Methylpiperazin-1-ylmethyl)-N-[3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
 - Compound 7: Thiophene-3-carboxylic acid [4-methyl-3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-amide;
 - Compound 8: 4-Chloro-N-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-
20 phenyl]-benzamide;
 - Compound 9: 4-Chloro-N-[3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
 - Compound 10: 3,4,5-Trimethoxy-N-[3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
 - Compound 11: 4-Cyano-N-[3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-
30 benzamide;
 - Compound 12: 4-Methoxy-N-[3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
 - Compound 13: 4-Chloro-N-[3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-
35 benzenesulfonamide;
 - Compound 14: Thiophene-3-carboxylic acid [3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-amide;

- Compound 15: 3,5-Dimethoxy-*N*-[3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- Compound 16: 3,4,5-Trimethoxy-*N*-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 5 Compound 17: 4-Cyano-*N*-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- Compound 18: 4-Methoxy-*N*-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- Compound 19: 4-Chloro-*N*-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzenesulfonamide;
- 10 Compound 20: Thiophene-3-carboxylic acid [4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- Compound 21: 3,5-Dimethoxy-*N*-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 15 Compound 22: 4-Trifluoromethoxy-*N*-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- Compound 23: Cyclohexanecarboxylic acid [4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- Compound 24: Cyclohexanecarboxylic acid [3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- 20 Compound 25: Isoquinoline-5-sulfonic acid [4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- Compound 26: Isoquinoline-5-sulfonic acid [3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- 25 Compound 27: (5-Nitro-2-methylphenyl)-(4-pyridin-2-yl-pyrimidin-2-yl)-amine;
- Compound 28: (5-Amino-2-methylphenyl)-(4-pyridin-2-yl-pyrimidin-2-yl)-amine;
- Compound 29: 3,4,5-Trimethoxy-*N*-[4-methyl-3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 30 Compound 30: 4-Cyano-*N*-[4-methyl-3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- Compound 31: (3-Aminophenyl)-(4-pyridin-2-yl-pyrimidin-2-yl)-amine;
- Compound 32: 4-Chloro-*N*-[4-methyl-3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 35 Compound 33: Cyclohexanecarboxylic acid [4-methyl-3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- Compound 34: 4-Cyano-*N*-[4-methyl-3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;

- Compound 35: 4-Chloro-*N*-[4-methyl-3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzenesulfonamide;
- Compound 36: 4-Methoxy-*N*-[4-methyl-3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 5 Compound 37: 4-Chloro-*N*-[4-methyl-3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- Compound 38: Cyclohexanecarboxylic acid [3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- Compound 39: 3,5-Dimethoxy-*N*-[3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 10 Compound 40: (5-Amino-2-methylphenyl)-(4-pyridin-4-yl-pyrimidin-2-yl)-amine;
- Compound 41: Thiophene-3-carboxylic acid [3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- 15 Compound 42: 4-Chloro-*N*-[3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzenesulfonamide;
- Compound 43: 4-Chloro-*N*-[3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- Compound 44: (3-Aminophenyl)-(4-pyridin-4-yl-pyrimidin-2-yl)-amine;
- 20 Compound 45: (3-Nitrophenyl)-(4-pyridin-4-yl-pyrimidin-2-yl)-amine;
- Compound 46: 4-Trifluoromethoxy-*N*-[4-methyl-3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- Compound 47: Isoquinoline-5-sulfonic acid [3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- 25 Compound 48: 4-Methoxy-*N*-[3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- Compound 49: 4-Cyano-*N*-[3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- Compound 50: 3,4,5-Trimethoxy-*N*-[3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 30 Compound 51: 3,5-Dimethoxy-*N*-[4-methyl-3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- Compound 52: 3,4,5-Trimethoxy-*N*-[4-methyl-3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 35 Compound 53: 4-(4-Methylpiperazin-1-ylmethyl)-*N*-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide (GleevecTM);
- Compound 54: 4-Methyl-*N*-[4-methyl-3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzenesulfonamide

- Compound 55: 4-Methoxy-*N*-[4-methyl-3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- Compound 56: 3,5-Dimethoxy-*N*-[4-methyl-3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 5 Compound 57: Naphthalene-2-carboxylic acid [4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- Compound 58: *N*-[3-(4-Pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- Compound 59: 4-Chloro-*N*-[3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 10 Compound 60: 4-Methoxy-*N*-[3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- Compound 61: 4-Chloro-*N*-[3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzenesulfonamide;
- Compound 62: Thiophene-2-carboxylic acid 3-(4-pyridin-2-yl-pyrimidin-2-yl-15 amino)-phenyl]-amide;
- Compound 63: Naphthalene-2-sulfonic-acid [3-(4-pyridin-2-yl-pyrimidin-2-yl-amino)-phenyl]-amide;
- Compound 64: Isoquinoline-5-sulfonic-acid [3-(4-pyridin-2-yl-pyrimidin-2-yl-amino)-phenyl]-amide;
- 20 Compound 65: Cyclopentanecarboxylic acid 3-(4-pyridin-2-yl-pyrimidin-2-yl-amino)-phenyl]-amide;
- Compound 66: Naphthalene-2-carboxylic acid [3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- Compound 67: 4-Cyano-*N*-[3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-25 benzamide;
- Compound 68: 3,5-Dimethoxy-*N*-[3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- Compound 69: 4-Bromo-*N*-[3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 30 Compound 70: 4-Methyl-*N*-[3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- Compound 71: 4-Fluoro-*N*-3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzenesulfonamide;
- Compound 72: 3,5-Dichloro-*N*-[3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-35 benzamide;
- Compound 73: *N*-[3-(4-Pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- Compound 74: 4-Chloromethyl-*N*-[3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;

- Compound 75: 4-Methyl-*N*-3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzenesulfonamide
- Compound 76: 4-(4-Methylpiperazin-1-ylmethyl)-*N*-[3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 5 Compound 77: Naphthalene-2-carboxylic acid [3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- Compound 78: 2-Methoxy-*N*-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- Compound 79: 2-Methoxy-*N*-[3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 10 Compound 80: 4-Methyl-*N*-[3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- Compound 81: 4-Methyl-*N*-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 15 Compound 82: *N*-[4-Methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- Compound 83: 1-(3,5-Diacetyl-phenyl)-3-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-urea;
- Compound 84: 1-{3,5-Bis-(amidinohydrazone)-phenyl}-3-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-urea;
- 20 Compound 85: *N*-[4-Methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-nicotinamide;
- Compound 86: *N*-[3-(4-Pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-nicotinamide;
- Compound 87: [1,8]Naphthyridine-2-carboxylic acid [3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- 25 Compound 88: [1,8]Naphthyridine-2-carbothioic acid [3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- Compound 89: 2-Methoxy-*N*-[3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 30 Compound 90: 4-Trifluoromethoxy-*N*-[3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- Compound 91: 4-Methyl-*N*-[3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 35 and pharmaceutically active salts of these compounds.

Recent research has revealed how cells communicate with each other to coordinate the growth and maintenance of the multitude of tissues within the

human body. A key element of this communication network is the transmission of a signal from the exterior of a cell to its nucleus, which results in the activation or suppression of specific genes. This process is called signal transduction.

- 5 An integral part of signal transduction is the interaction of ligands, their receptors and intracellular signal transduction molecules. Ligands are messengers that bind to specific receptors on the surface of target cells. As a result of the binding, the receptors trigger the activation of a cascade of downstream signaling molecules, thereby transmitting the message from the exterior of the cell to its
- 10 nucleus. When the message reaches the nucleus, it initiates the modulation of specific genes, resulting in the production of RNA and finally proteins that carry out a specific biological function. Disturbed activity of signal transduction molecules may lead to the malfunctioning of cells and disease processes. Specifically, interference of the pathogenic PrP^{Sc} from prion diseases with
- 15 neuronal cells is necessary for the prion protein to induce its neuropathological features such as neuronal vacuolization, neuronal death and gliosis with hyperastrocytosis.

A key element of this communication network is the transmission of a signal from

20 the exterior of a cell to its nucleus, which results in the activation or suppression of specific genes. The human cellular protein kinases Abl and clk1 are two of the enzymes involved in said signal transduction process. As revealed herein said kinases Abl and clk1 serve as targets and are inhibited by the pyridylpyrimidine compounds of the general formula (I). It could be proved that prion infections

25 and/or prion diseases can be treated and also be prevented by the inhibition of said kinase Abl using the inventive pyridylpyrimidine derivatives. Inhibition of the kinase clk1 by said pyridylpyrimidine compounds can be used for the treatment of infections and diseases.

- 30 A microarray platform technology consisting of more than 1100 signal transduction cDNAs has been established. The technology is used for the identification of changes in RNA expression patterns as a result of the manipulation of the host cell by PrP^{Sc}. In addition, differential display techniques were used in order to pinpoint these changes to those enzymes which could be potential targets for drug
- 35 intervention.

Employing this predefined set of signal transduction relevant cDNAs on the filters, the expression pattern of signal transduction mRNAs in neuronal mouse cells

transfected with the pathogenic form of the prion protein (PrP^{Sc}) were compared with the same cells transfected with the non-pathogenic wild-type form (PrP^C) as a control. Interference of the PrP^{Sc} with the cellular signaling events is reflected in different gene expression when compared to the control cellular situation (PrP^C).

5

Using this technology, the human cellular protein kinases FGF-R1 (also known as flg, Flt-1, Flt-2, or b-FGFR), Tkt (also known as CCK-2, DDR-2, or EDDR, EC Number 2.7.1.112), Abl (also known as c-abl), clk1, MKK7 (also known as SKK4, SAPKK4, SAPKK5, or JNKK2), LIMK-2, CaM-KI, JNK2 (also known as SAPK1a, SAPKalpha), CDC2 (also known as CDK1), PRK, the human cellular protein phosphatases PTP-SL (also known as MCP83), PTP-zeta, the cellular signal transduction molecules HSP86, and GPIR-1 were identified as potential anti-prion disease targets. Said cellular protein kinases, phosphatases and signal transduction molecules are found to be specifically up- or downregulated by PrP^{Sc} in relevant mouse neuronal cells.

15

Surprisingly, it was found that the following human cellular targets are significantly up- or downregulated in prion infected cells:

20

target	regulation
FGF-R1	3.6 fold stronger
Abl	5.6 fold stronger
MKK7	4.1 fold stronger
CDC2	2.0 fold weaker
25	Tkt
	2.1 fold stronger
	LIMK-2
	2.1 fold stronger
	CaM-KI
	2.1 fold stronger
	JNK2
	2.0 fold weaker
	PRK
	2.0 fold weaker
30	PTPzeta
	4.6 fold weaker
	PTP-SL
	5.0 fold weaker
	HSP86
	4.1 fold weaker
	GPIR-1
	2.3 fold weaker

35

Thus, one aspect of the present invention relates to a method for preventing and/or treating prion infections and/or diseases associated with said prion infections in an individual which comprises administering to the individual an amount of at least one compound of the general formula (I) and/or pharmaceutically acceptable salts

thereof effective to prevent and/or treat said prion infections and/or prion diseases. Most preferred is the administration of a compound according to claim 8.

It could be proven that inhibition of one target selected from FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1 was effective to treat prion diseases. Therefore, another aspect of the invention relates to a method for preventing and/or treating prion infections and/or prion diseases in an individual comprising the step of administering a pharmaceutically effective amount of at least one compound according of the general formula (I) and/or pharmaceutically acceptable salts thereof which inhibits at least partially the activity of one target selectef from FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1.

The nucleoside sequences of the genes coding for the human cellular protein kinase Abl and the protein kinase clk1 and their amino acid sequences are disclosed in form of a sequence listing shown below. The nucleoside and amino acid sequences for the kinase Abl (Accession Number: M14752) and for the kinase clk1 (Accession Numbers: XM002520, NM004071, L29222, L29219) were obtained from NCBI (National Library of Medicine: PubMed).

The compounds of general formula (I) were identified as inhibitors of at least one target selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, GPIR-1 by the use of a method for detecting compounds useful for the prophylaxis and/or treatment of prion infections and/or diseases. Said method comprises

- a) contacting a test compound with at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, GPIR-1; and
- b) detecting the activity of said human cellular protein kinase, phosphatase or cellular signal transduction molecule.

The activity of a human cellular protein kinase, phosphatase or cellular signal transduction molecule was preferably measured by means of an enzymatic assay.

As used herein, the term "inhibitor" refers to any compound capable of downregulating, decreasing, suppressing or otherwise regulating the amount and/or activity of at least one human cellular protein kinase, phosphatase or cellular

signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1. Generally, said inhibitors, including suicide inhibitors, may be proteins, oligo- and polypeptides, nucleic acids, genes, small chemical molecules, or other chemical moieties.

The present disclosure teaches for the first time the up- or downregulation of the above-mentioned human cellular protein kinases, phosphatases, or cellular signal transduction molecules specifically involved in prion infections and/or diseases.

Thus, the present invention is also directed to a method for detecting prion infections and/or diseases in an individual comprising:

- a) providing a sample from said individual; and
- b) adding to said sample a pharmaceutically effective amount of at least one pharmaceutically active agent; and
- c) detecting activity in said sample of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, GPIR-1.

As used herein the term "sample" refers to any sample that can be taken from a living animal or human for diagnostic purposes, especially said sample comprises blood, milk, saliva, sputum, excrement, urine, spinal cord liquid, liquor, lachrymal gland liquid, biopsies and all other samples that can be taken from a living animal or human for diagnostic purposes.

The term "individual" preferably refers to mammals, especially humans or ruminants. Ruminants are, for instance, muledeer, elk, cow, cattle, sheep, goat, deer, or buffalo. Minks are an example for mammals which do not belong to the species of ruminants.

As used herein the term "ruminants" refers to an animal, for instance, cattle, sheep, goat, deer, elk, or buffalo that has four separate stomach chambers, and is therefore able to digest a wide range of organic and plant foods. The term "ruminants" refers also to exotic ruminants, like captive nyala, gemsbok, Arabian oryx, eland, kudu, scimitar-horned oryx, ankole, or bison which are also accessible to develop spongiform encephalopathy.

A similar aspect of the present invention is directed to a method for detecting prion infections and/or prion diseases in cells, cell cultures and/or cell lysates comprising:

- a) providing said cells, cell cultures and/or cell lysates; and
- b) adding to said cells, cell cultures and/or cell lysates a pharmaceutically effective amount of at least one pharmaceutically active agent; and
- c) detecting activity in said sample of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, GPIR-1.

Furthermore, it has been shown that the inhibition of at least one target selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, GPIR-1 has an effect on the production of prions. Therefore, another aspect of the invention relates to a method for regulating the production of prions in an individual or in cells comprising the step of administering a pharmaceutically effective amount of at least one pharmaceutically active agent which inhibits at least partially the activity of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, GPIR-1, or which inhibits at least partially the production of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, GPIR-1.

The inventive compounds according to general formula (I) are examples for the above-mentioned pharmaceutically active agent. Preferably the targets FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, and CDC 2 are used with said methods.

Another type of pharmaceutically active agents useful within the methods disclosed herein are monoclonal or polyclonal antibodies which bind to a human cellular protein kinase, phosphatase or a cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1. Thus, a further aspect of the present invention is related to said monoclonal or polyclonal antibodies which bind to a human cellular protein kinase, phosphatase or a cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1,

MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1.

Another embodiment of the present invention utilizes the scientific findings that
5 some targets such as JNK2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1 are
downregulated during prion infection and that upregulation of the effected target
by means of an activator leads to an alternative way of treating prion infections
and diseases associated with prion infection.

10 Thus, a method was developed for regulating the production of prions either in an
individual or in cells. Said methods comprise the step of administering an individual
or the cells a pharmaceutically effective amount of at least one pharmaceutically
active agent wherein said agent activates at least partially the activity of at least one
15 human cellular protein kinase, phosphatase or cellular signal transduction
molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7,
LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1, or
wherein said agent at least partially activates or stimulates the production of at
least one human cellular protein kinase, phosphatase or cellular signal
20 transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1,
MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and
GPIR-1.

Preferably the targets JNK2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1 are
used within the above-described methods.

25 Because of the fact that the organism may upregulate a given target such as FGF-
R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, and CDC 2 in order to compete with
the prion infection, it is also a reasonable approach to further support said
upregulation by means of an activator. Therefore, the above-mentioned methods
30 apply either to targets which are downregulated but also to targets which are
upregulated.

The novel and partially known pyridylpyrimidine compounds of the general formula
(I) represent a new class of pharmaceuticals highly useful for the prophylaxis and
35 treatment of prion infections and prion diseases.

Thus, a further aspect of the present invention describes the use of a compound of
the general formula (I) and/or pharmaceutically acceptable salts thereof for the

manufacture of a pharmaceutical formulation for prophylaxis and/or treatment of prion infections and/or diseases induced or caused by prion infection.

5 As used herein the Term "prion diseases" refers to transmissible spongiform encephalopathies. This group of neurologic diseases affects humans and many species of animals causing a "sponge-like" degeneration of brain tissue. Among other unique features, all of these diseases are associated with the accumulation of an abnormal form of the prion protein in nerve cells that eventually leads to the death of the host. While prion diseases can all be transmitted from one host to another, it remains contentious as to whether a virus-like infectious agent or the abnormal prion protein itself, the prion, causes the conversion of normal to abnormal protein.

15 Probably most mammalian species develop prion diseases. Specific examples for animals include:

- **Scrapie** sheep, goat
- **TME** (transmissible mink encephalopathy): mink
- **CWD** (chronic wasting disease): muledeer, deer, elk
- **BSE** (bovine spongiform encephalopathy): cows, cattles

20

Humans are also susceptible to several prion diseases. Examples are:

- **CJD** Creutzfeld-Jacob Disease
- **GSS** Gerstmann-Sträussler-Scheinker syndrome
- **FFI** Fatal familial Insomnia
- 25 • **Kuru**
- **Alpers Syndrome**

The human prion diseases include kuru, sporadic Creutzfeldt-Jakob disease (sCJD), familial CJD (fCJD), iatrogenic CJD (iCJD), Gerstmann-Sträussler-Scheinker (GSS) disease, fatal familial insomnia (FFI), and, more recently, new variant CJD (nvCJD or vCJD). In addition to these human diseases, prion-related

30

diseases, have been recognized in several animal hosts. Scrapie is a naturally occurring disease of sheep and goats that causes ataxia, behavioral changes, and a severe pruritus that leads to scraping behavior, from which the disease was named. Additional prion diseases in animals include transmissible mink encephalopathy (TME), chronic wasting disease (CWD) of deer and elk, feline spongiform encephalopathy (FSE), and bovine spongiform encephalopathy (BSE), among others.

The transmissible nature of prion disease was first demonstrated experimentally in 1936 when Cuillé and Chelle transmitted scrapie to a healthy goat by the intraocular administration of scrapie-infected spinal cord. Thirty years later, sCJD was transmitted to chimpanzees. The pathologic feature common to all these diseases is a prominent vacuolation of the gray matter of the brain that produces a "sponge-like" appearance on light microscopy. This histopathologic appearance, coupled with the transmissible nature of these diseases, led to their collective designation as "transmissible spongiform encephalopathies" or TSEs.

The etiologic agent of the TSEs was proposed to be a "slow virus" to explain its transmissible nature and the prolonged incubation period observed during experimental transmission studies. Early experiments suggested that protein may be a critical component of the infectious agent. These studies established the basis for a new form of a transmissible pathogen, one that is composed ostensibly of only protein and lacks any replicative elements such as nucleic acid.

The term "prion" was coined to indicate an *infectious* agent with *proteinlike* properties. The unusual properties of the pathogen were demonstrated in early experiments in which conditions that degrade nucleic acids, such as exposure to ionizing and ultraviolet radiation, did not reduce the infectivity of scrapie fractions. On the other hand, treatments that degrade protein, such as prolonged exposure to proteases, correlated with a reduction in infectivity. A protein with relative resistance to protease digestion was found to be consistently present in the brains of animals and humans with TSE. Surprisingly, this protein was found to be one that is normally encoded by a chromosomal gene of the host.

Thus, the question raised, how a normally expressed protein could also be a transmissible pathogen? It was hypothesized and later demonstrated that PrP exists in two major isoforms: the nonpathogenic or cellular form, designated PrP^C, and the pathogenic or scrapie-inducing form, designated PrP^{Sc}. Both PrP^C and

PrP^{Sc} have the same amino acid sequence, yet they differ in their biochemical properties: PrP^C is soluble in nondenaturing detergents and completely degraded by proteases, whereas PrP^{Sc} is insoluble in nondenaturing detergents and shows a relative resistance to proteases. Structural studies of PrP^C and PrP^{Sc} indicate a
5 difference in the conformation of the two isoforms: PrP^C is predominantly helical, whereas PrP^{Sc} contains at least 40% pleated sheet structure. Conversion to this sheet structure appears to be the fundamental event in prion disease. The ultimate mechanism of how cells die coincident with the generation of prions is still unclear. Simple accumulation of pathogenic protein may not be sufficient to
10 explain disease, however, it may constitute a critical step in cellular dysfunction.

It was shown that the pyridylpyrimidine compounds of the general formula (I) are highly effective for the prophylaxis and/or treatment of prion infections and/or prion diseases selected from the group comprising Scrapie, TME, CWD, BSE, CJD,
15 vCJD, GSS, FFI, Kuru, and Alpers Syndrome. Preferably, the pyridylpyrimidine derivatives are used for preventing and/or treating BSE, vCJD, or CJD.

The above-mentioned prion infections and/or diseases associated with prion infections can be treated using the inventive pyridylpyrimidine derivatives by
20 targeting at least one of the human cellular protein kinases, phosphatases or cellular signal transduction molecules selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, GPIR-1. Thereby, the compounds according to general formula (I) act as inhibitors for at least one of the above-mentioned targets and especially as
25 inhibitors for at least one enzyme selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, and CDC 2.

According to these findings a further aspect of the present invention is directed to a method for preventing and/or treating prion infections and/or prion diseases in an
30 individual comprising the step of administering a pharmaceutically effective amount of at least one pharmaceutically active agent which inhibits at least partially the activity of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86,
35 GPIR-1, or which inhibits at least partially the production of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, GPIR-1.

Another aspect is related to a method for preventing and/or treating prion infections and/or prion diseases in cells or cell cultures comprising the step of administering a pharmaceutically effective amount of at least one pharmaceutically active agent which inhibits at least partially the activity of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, GPIR-1, or which inhibits at least partially the production of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, GPIR-1.

The inventive pyridylpyrimidine compounds of formula (I) are examples for the above-mentioned inhibitor. Said pyridylpyrimidine compounds and/or pharmaceutically acceptable salts thereof are administered in a dosage corresponding to an effective concentration in the range of 0.01 – 50 μ M, preferably in the range of 0.01 – 10 μ M, more preferably in the range of 0.01 – 1 μ M, and most preferably in the range of 0.01 – 0.1 μ M.

Because of the fact that the targets JNK2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1 are downregulated in cells infected with prions, an upregulation of said targets represents another strategy in order to treat prion infections and diseases like CJD (nvCJD or vCJD) associated with prion infections. Said upregulation can be performed by activators.

An agent that is able to upregulate, increase, activate, or stimulate the activity of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1, but especially of JNK2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1 is named "activator".

Thus, another embodiment of the present invention describes a method for preventing and/or treating prion infections and/or diseases in an individual comprising the step of administering a pharmaceutically effective amount of at least one pharmaceutically active agent which activates at least partially the activity of at least one human cellular protein kinase, phosphatase or cellular signal

transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1, or which activates or stimulates the production of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule
5 selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1. Preferably, said method is directed to the targets JNK2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1.

- 10 As used herein, the term "agent" or "pharmaceutically active agent" refers to any chemical compound capable of down- or upregulating, de- or increasing, suppressing, activation, stimulating or otherwise regulating the amount and/or activity of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt,
15 Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1. Generally, said agents may be proteins, oligo- and polypeptides, nucleic acids, genes, aptamers, small chemical molecules, or other chemical moieties. An agent may be either an inhibitor or an activator and especially an inhibitor for the enzymes FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, and
20 CDC 2 and an activator for the targets JNK2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1.

One special kind of said pharmaceutically active agents are aptamers which function as regulators of the activity of a wide range of cellular molecules such as
25 human cellular protein kinase and phosphatase. Aptamers are nucleic acid molecules selected in vitro to bind small molecules, peptides, or proteins with high affinity and specificity. Aptamers not only exhibit highly specific molecular recognition properties but are also able to modulate the function of their cognate targets in a highly specific manner by agonistic or antagonistic mechanisms.
30 Most famous examples for aptamers are DNA aptamers or RNA aptamers.

Further examples for pharmaceutically active agents are the pyridylpyrimidine compounds of the present invention and/or pharmaceutically acceptable salts thereof. Said compounds are administered in a dosage corresponding to an
35 effective concentration in the range of 0.01 – 50 μ M, preferably in the range of 0.01 – 10 μ M, more preferably in the range of 0.01 – 1 μ M, and most preferably in the range of 0.01 – 0.1 μ M.

The compounds of general formula (I) can be administered in a daily dosage in the range of 25 mg to 1000 mg, preferably in a daily dosage of 400 mg to 600 mg, more preferably in a daily dosage of 500 mg, and most preferably in continuously increased daily dosages starting at a initial daily dosage of 400 mg and ending up in a daily dosage of 600 mg at the end of the treatment.

A question is how PrP^C does convert to PrP^{Sc}? Potential mechanisms that initiate conversion of PrP^C to PrP^{Sc} include a germ line mutation of the human prion protein gene (PRNP), a somatic mutation within a particular neuron, and spontaneous conversion of PrP^C to an aberrant conformation that is not refolded appropriately to its native structure. The prion protein gene (PRNP) is the single gene on the short arm of chromosome 20 in humans which encodes the normal cellular isoform of the prion protein. Regardless of the initiating event, once an "infectious unit" has been generated, PrP^{Sc} appears to act as a conformational template by which PrP^C is converted to a new molecule of PrP^{Sc} through protein-protein interaction of PrP^{Sc} and PrP^C. This concept is supported by several studies which show that mice with the normal PrP gene deleted (PrP knockout mice) do not develop prion disease after inoculation with scrapie. Furthermore, transgenic (Tg) mice that express a chimeric PrP gene made of human and mouse segments develop protease-resistant chimeric mouse-human PrP^{Sc} in their brains when inoculated with brain extracts from humans with prion disease. These findings clearly illustrate that prions do not self-replicate but instead convert nonpathogenic PrP^C to pathogenic PrP^{Sc}.

In its sporadic or nonfamilial form, **CJD** is the most common of the human prion diseases. Confusion and forgetfulness which progress rapidly to severe cortical dementia in combination with ataxia, myoclonus, and an abnormal electroencephalogram (EEG) represents the "classic tetrad" of CJD. However, a host of other neurologic signs and symptoms, including diffuse or focal weakness, painful neuropathy, choreiform movements, hallucinations, cortical blindness, primary language disturbance, supranuclear ophthalmoplegia, and alien hand syndrome, among others, have been observed. As the disease progresses from the early stage, ataxia commonly limits the patient's mobility.

Familial CJD (fCJD) includes those cases with a dominantly inherited mutation of the *PRNP* gene, in which the pathologic features of spongiform change occur in the absence of GSS-type plaques. Although, familial cases of CJD tend to have a clinical and pathologic phenotype similar to that of sCJD.

The original description of a patient with the onset of ataxia and dysarthria followed by variable degrees of pyramidal and extrapyramidal symptoms and late developing dementia defines the classic presentation of **GSS**. The duration of said disease ranges from 2 to 10 years. Death usually results from secondary infection, often from aspiration pneumonia because of impaired swallowing. The presence of plaque deposits regionally or diffusely throughout the cortex that are immunoreactive to anti-human PrP antibodies is the hallmark of this form of prion disease.

10

FFI is a genetic disorder which manifests itself by many symptoms due to the degeneration of a certain part of the brain, the thalamus. The affected area of the brain is the area responsible for sleep, the thalamus. The thalamus is the center which communications from the brain to the body and the body to the brain pass through for proper directions to where a signal should be received. When sleep takes place, it is thought that the thalamus becomes less efficient at this signal transfer function allowing for the vegetative state of sleep to come over an individual. Consequently, the symptoms of fatal familial insomnia are directly related to the malfunction of the responsibilities of the thalamus, namely sleep.

20

There are four stages of the disease before an individual's life ends. The first stage is progressive insomnia, the characteristic feature of fatal familial insomnia. By now, there is no cure for this illness.

25 The term "familial" means: affecting several members of the same family, usually as a result of an underlying genetic mutation.

The occurrence of **vCJD** is sobering because it appears to represent a situation in which the prion has "jumped" species, in this case from cow to human. Because the pathologic features and clinical presentation of vCJD differ significantly from those of sCJD, it is considered a new "strain" of human prion disease. The same "protein signature" was observed following experimental transmission of BSE to several animal hosts, supporting the idea that vCJD results from the infection of humans with BSE. vCJD occurs primarily in younger individuals (average age 27) with a somewhat protracted course of approximately 16 months. The brain shows diffuse vacuolation and the presence of distinctive dense core PrP-containing plaques surrounded by a halo of spongiform change.

30
35

Kuru is the condition which first brought prion diseases to prominence in the 1950s. The disease was found in geographically isolated tribes in New Guinea. It was established that ingesting brain tissue of dead relatives for religious reasons was likely to be the route of transmission.

5

Alpers Syndrome is the name given to prion diseases in infants.

Scrapie is the accepted, albeit somewhat colloquial, name for the naturally occurring transmissible spongiform encephalopathy of sheep and goats found worldwide. Scrapie also infects laboratory mice and hamsters making it one of the most important sources of new scientific information about this group of disorders. Scrapie was the first example of this type of disease to be noticed and has been known about for many hundreds of years. There are two possible methods of transmission in sheep: a) Infection of pasture with placental tissue carrying the agent followed by ingestion, or b) direct sheep-lamb transmission.

15

CWD is a fatal neurodegenerative disease of deer and elk, now known to be a transmissible spongiform encephalopathy. To date, affected animals have been found exclusively in the United States.

20

BSE

Bovine spongiform encephalopathy or "mad cow disease" appears to have originated from scrapie that has been recognized in Europe since the mid-18th century. It has since spread to most sheep-breeding countries and is widespread in the United Kingdom, where until 1988 the rendered carcasses of livestock (including sheep) were fed to ruminants and other animals as a protein-rich nutritional supplement.

25

During rendering, carcasses from which all consumable parts had been removed were milled and then decomposed in large vats by boiling at atmospheric or higher pressures, producing an aqueous slurry of protein under a layer of fat (tallow). After the fat was removed, the slurry was desiccated into a meat and bone meal product that was packaged by the animal food industry and distributed to owners of livestock and other captive animals (e.g., zoo and laboratory animals, breeding species, pets).

35

A further aspect is related to a method for regulating the expression of at least one human cellular protein kinase, phosphatase or cellular signal transduction

molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1 in an individual comprising the step of administering the individual a pharmaceutically effective amount of at least one pharmaceutically active agent wherein said agent
5 inhibits at least partially the transcription of DNA or the translation of RNA.

And a still further aspect of the present invention relates to a method for regulating the expression of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1,
10 Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1 in the cells, the method comprising the step of administering the cells a pharmaceutically effective amount of at least one pharmaceutically active agent wherein said agent inhibits at least partially the transcription of DNA or the translation of RNA.

15 As used herein, the term "regulating expression and/or activity" generally refers to any process that functions to control or modulate the quantity or activity (functionality) of a cellular component. Static regulation maintains expression and/or activity at some given level. Upregulation refers to a relative increase in
20 expression and/or activity. Accordingly downregulation refers to a relative decrease in expression and/or activity. Downregulation is synonymous with inhibition of a given cellular component's activity.

The transcription of DNA and the translation of RNA can be inhibited by
25 oligonucleotides or oligonucleotide derivatives. Thus, the present invention discloses oligonucleotides and derivatives of oligonucleotides which may be used in the above-mentioned methods. The oligonucleotide and/or its derivatives bind to the DNA and/or RNA encoding a human cellular protein kinase, phosphatase or a cellular signal transduction molecule selected from the group comprising FGF-R1,
30 Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1 and suppress the transcription of DNA or translation of RNA.

As described above, said prion infection and/or disease associated with said prion infection is selected from the group comprising Scrapie, TME, CWD, BSE, vCJD, CJD, GSS, FFI, Kuru, and Alpers Syndrome. Preferably, the method is used for prophylaxis and/or treatment of BSE, vCJD, or CJD. The above disclosed methods are preferably applied to CJD, vCJD, and BSE, more preferably applied to vCJD and BSE, and most preferably applied to BSE.
35

Some methods of the present invention identify compounds useful for prophylaxis and/or treatment of prion infections and/or diseases by screening a test compound, or a library of test compounds, for its ability to inhibit at least one of the above-mentioned human cellular protein kinases, phosphatases, or cellular signal transduction molecules, identified herein as characteristically up- or downregulated during prion production or growth inside a cell or individual. A variety of assay protocols and detection techniques are well known in the art and easily adapted for this purpose by a skilled practitioner. Such methods include, but are not limited to, high throughput assays (e.g., microarray technology, phage display technology), and *in vitro* and *in vivo* cellular and tissue assays.

Thus, a solid support is disclosed in the present invention useful for screening compounds useful for the prophylaxis and/or treatment of prion infections and/or diseases in an individual, the solid support comprising at least one immobilized oligonucleotide, wherein said oligonucleotide encodes one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1.

A further aspect of the present invention is related to a solid support useful for screening compounds useful for the prophylaxis and/or treatment of prion infections and/or diseases in an individual, the solid support comprising at least one immobilized human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1.

In another embodiment, a component of the above-mentioned methods comprises peptide fragments of one or more of the above-identified human cellular protein kinases, phosphatases or cellular signal transduction molecules immobilized on a solid support. Once again the most preferred solid support embodiment would contain polymers of sufficient quality and quantity to detect all of the above-mentioned human cellular protein kinases, phosphatase and cellular signal transduction molecules (e.g., a nucleic acid or a peptide microarray). A variety of supports and constructions of the same for the methods disclosed herein are well known in the art and easily adapted for this purpose by a skilled practitioner (cf.,

for example: Marschall, 1999 "Do-it-yourself gene watching" Science 286, 444-447; Service 2000 "Protein arrays step out of DNA's shadow" Science 289, 1673).

It is preferred that mRNA is measured as an indication of expression. Methods
 5 for assaying for mRNA include, but are not limited to, Northern blots, slot blots, dot blots, and hybridization to an ordered array of oligonucleotides. Nucleic acid probes useful for assay of a sample are preferably of sufficient length to specifically hybridize only to appropriate, complementary transcripts. Typically the oligonucleotide probes will be at least 10 to 25 nucleotides in length. In some
 10 cases longer probes of at least 30 to 50 nucleotides will be desirable.

The cDNA oligonucleotides immobilized on said membrane filter which are used for detecting the up- or downregulation of the above-mentioned human cellular protein kinases, phosphatases, and cellular signal transduction molecules by
 15 hybridization to the radioactively labeled cDNA probes have the nucleotide sequences listed in table 1.

Table 1: Nucleotide sequences of cDNA-arrays

Cellular kinase, phosphatase, or signal transduction molecule	Sequence of immobilized DNA on arrays (in relation to the respective Acc No)
FGF-R1	41 bp – 2619bp (X52833)
Tkt (EC 2.7.1.112)	1 bp – 3096bp (X74764)
Abl	2153 bp – 3765 bp (M14752)
clk1	156 bp -1610 bp (L29219)
MKK7	77 bp – 1323 bp (AF013588)
CDC2	77 bp – 1050 bp (X05360)
CaMKI	145 bp – 1452 bp (L41816)
JNK2	507 bp – 1782 bp (L31951)
LIMK-2	963 bp – 2047 bp (D45906)
PRK	n.a bp – 1862 bp (U56998)
PTP zeta (EC 3.1.3.48)	148 bp – 7604 bp (X54135)
PTP-SL	862 bp – 1902 bp (NM 002849)
HSP86	n.a bp – n.a bp (X07270)
GPIR-1	n.a bp – n.a bp (n.a)

20

Tkt has been assigned to the EC Number: 2.7.1.112

PTP zeta has been assigned to the EC Number : 3.1.3.48

The nucleoside sequences of the genes coding for the human cellular protein kinases, phosphatases, or cellular signal transduction molecules listed in Table 1 together with the amino acid sequences and the enzyme commission numbers (E.C. numbers) of said enzymes can be obtained from NCBI (National Library of Medicine: PubMed; Web address: www.ncbi.nlm.nih.gov/entrez).

The polypeptide product of gene expression may be assayed to determine the amount of expression as well. Methods for assaying for a protein include, but are not limited to, western blot, immuno-precipitation, radioimmuno assay, and peptide immobilization in an ordered array. It is understood, however, that any method for specifically and quantitatively measuring a specific protein or mRNA product can be used.

A variety of supports upon which nucleic acids or peptides can be immobilized are known in the art, for example filters, or polyvinyl chloride dishes. Any solid surface to which oligonucleotides or peptides can be bound, either directly or indirectly, either covalently or non-covalently, can be used. A preferred solid support is a microarray membrane filter or a "biochip". These contain particular polymer probes in predetermined locations on the array. Each predetermined location may contain more than one molecule of the probe, but each molecule within the predetermined location has an identical sequence.

The present invention incorporates by reference in their entirety techniques well known in the field of molecular biology. These techniques include, but are not limited to, techniques described in the following publications:

Ausubel, F.M. et al. eds., "Short Protocols In Molecular Biology" 4th Ed. 1999, John Wiley & Sons, NY (ISBN 0-471-32938-X);

Old, R.W. & S.B. Primrose "Principles of Gene Manipulation: An Introduction To Genetic Engineering" 3rd Ed. 1985, Blackwell Scientific Publications, Boston. Studies in Microbiology: V.2, 409 pp. (ISBN 0-632-01318-4);

Mayer, R.J. & J.H. Walker eds. "Immunochemical Methods In Cell and Molecular Biology" 1987, Academic Press, London. 325 pp. (ISBN 0-12480-855-7);

Winnacker, E.L. "From Genes To Clones: Introduction To Gene Technology" 1987 VCH Publishers, NY. (translated by Horst Ibelgaufts) 634 pp. (ISBN 0-89573-614-4).

As described above, a microarray platform technology was developed consisting of more than 1100 signal transduction cDNAs immobilized on a solid support. Thus, another aspect of the present invention is directed to a solid support useful for detecting prion infections and/or diseases in an individual, the solid support comprising an immobilized oligonucleotide, wherein said oligonucleotide is capable of detecting activity of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1.

The present invention discloses also for the first time a solid support useful for detecting prion infections and/or diseases in cells, the solid support comprising an immobilized oligonucleotide, wherein said oligonucleotide is capable of detecting activity of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1.

The present invention further incorporates by reference in their entirety techniques well known in the field of microarray construction and analysis. These techniques include, but are not limited to, techniques described in the following patents and patent applications describing array of biopolymeric compounds and methods for their fabrication:

U.S. Pat. Nos. 5,807,522; 6,087,102; WO 93/17126; WO 95/11995; WO 95/35505; EP 742 287; and EP 799 897.

Techniques also include, but are not limited to, techniques described in the following patents and patent application describing methods of using arrays in various applications:

U.S. Pat. Nos. 5,994,076; 6,033,860; 6,040,138; 6,040,140; WO 95/21265; WO 96/31622; WO 97/10365; WO 97/27317; EP 373 203; and EP 785 280

Still a further aspect of the present invention is directed to pharmaceutical compositions comprising at least one pharmaceutically active agent together with a pharmaceutically acceptable carrier, excipient or diluents. Examples for

pharmaceutically active agents are the above-mentioned inventive compounds according to formula (I), or other small chemical molecules, antibodies, aptamers, oligo- and polynucleotides, genes and other biological components capable of regulating the activity of at least one target selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1, or which are effective to treat prion infections and diseases associated with prion infection. Said prion infections and diseases are preferably Scrapie, TME, CWD, BSE, vCJD, CJD, GSS, FFI, Kuru, and Alpers Syndrome.

10

Thus, the pharmaceutical compositions according to the present invention may comprise an inhibitor, such as the inventive pyridylpyrimidine compounds or an activator such as aptamers for at least one target selected from FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1. It is also possible to have a combination of inhibitors or activators as active ingredients in one single pharmaceutical composition. Furthermore, suitable are also combinations of at least one inhibitor and at least one activator for different targets within a single pharmaceutical composition. For example, a pharmaceutical composition could comprise compound 12 as an inhibitor for, for instance, the target Abl, and an activator such as an aptamer for, for instance, the human cellular protein kinase JNK2.

20

Said pharmaceutical compositions are useful for the prophylaxis and/or treatment of an individual afflicted with prions comprising at least one agent capable of inhibiting and/or activating at least partially the activity, the expression, and/or the production of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1.

25

30

The pyridylpyrimidine compounds of the present invention are basic and form pharmaceutically acceptable salts with organic and inorganic acids. Examples of suitable acids for such acid addition salt formation are hydrochloric acid, hydrobromic acid, sulfuric acid, phosphoric acid, acetic acid, citric acid, oxalic acid, malonic acid, salicylic acid, p-aminosalicylic acid, malic acid, fumaric acid, succinic acid, ascorbic acid, maleic acid, sulfonic acid, phosphonic acid, perchloric acid, nitric acid, formic acid, propionic acid, gluconic acid, lactic acid, tartaric acid, hydroxymaleic acid, pyruvic acid, phenylacetic acid, benzoic acid, p-aminobenzoic

35

acid, p-hydroxybenzoic acid, methanesulfonic acid, ethanesulfonic acid, nitrous acid, hydroxyethanesulfonic acid, ethylenesulfonic acid, p-toluenesulfonic acid, naphthylsulfonic acid, sulfanilic acid, camphorsulfonic acid, china acid, mandelic acid, o-methylmandelic acid, hydrogen-benzenesulfonic acid, picric acid, adipic acid, d-o-tolytartaric acid, tartronic acid, α -toluic acid, (o, m, p)-toluic acid, naphthylamine sulfonic acid, and other mineral or carboxylic acids well known to those skilled in the art. The salts are prepared by contacting the free base form with a sufficient amount of the desired acid to produce a salt in the conventional manner.

10

It is also possible to obtain acid addition salts with amino acids like methionine, tryptophane, lysine or arginine, especially with pyridylpyrimidine compounds of the general formula (I) carrying a carboxylic acid residue.

15 Depending upon the substituents on the inventive pyridylpyrimidine compounds, one may be able to form salts with bases, too. Thus, for example, if there are carboxylic acid substituents in the molecule, salts may be formed with inorganic as well as organic bases such as, for example, NaOH, KOH, NH₄OH, tetraalkylammonium hydroxide, and the like.

20

The compounds of the general formula (I) can also be administered in form of their pharmaceutically active salts optionally using substantially nontoxic pharmaceutically acceptable carriers, excipients or diluents. The medications of the present invention are prepared in a conventional solid or liquid carrier or diluents and a conventional pharmaceutically-made adjuvant at suitable dosage level in a known way. The preferred preparations are in administratable form which is suitable for oral application. These administratable forms, for example, include pills, tablets, film tablets, coated tablets, capsules, powders and deposits.

25

30 The preferred administratable forms are tablets, film tablets, coated tablets, gelatin capsules, and opaque capsules. Each pharmaceutical composition contains at least one compound of the general formula (I), preferably compound 53 and/or pharmaceutically acceptable salts thereof in an amount of 50 mg to 150 mg, preferably 80 mg to 120 mg, and most preferably in an amount of 100 mg per formulation.

35

Furthermor , the subject of the present invention also includes pharmaceutical preparations for parenteral, including dermal, intradermal, intragastrical,

intracutaneous, intravasal, intravenous, intramuscular, intraperitoneal, intranasal, intravaginal, intrabuccal, percutaneous, rectal, subcutaneous, sublingual, topical or transdermal application, which in addition to typical vehicles and diluents contain a pyridylpyrimidine compound of the general formula (I) and/or a
5 pharmaceutically acceptable salt thereof as active ingredient.

Within the disclosed methods the pharmaceutical compositions of the present invention, containing pyridylpyrimidine derivatives of the general formula (I) as active ingredients, will typically be administered in admixture with suitable carrier
10 materials selected with respect to the intended form of administration, i.e. oral tablets, capsules (either solid-filled, semi-solid filled or liquid filled), powders for constitution, oral gels, elixirs, dispersible granules, syrups, suspensions, and the like, and consistent with conventional pharmaceutical practices. For example, for oral administration in the form of tablets or capsules, the active drug component
15 may be combined with any oral nontoxic pharmaceutically acceptable inert carrier, such as lactose, starch, sucrose, cellulose, magnesium stearate, dicalcium phosphate, calcium sulfate, talc, mannitol, ethyl alcohol (liquid forms) and the like. Moreover, when desired or needed, suitable binders, lubricants, disintegrating agents and coloring agents may also be incorporated in the mixture. Powders and
20 tablets may be comprised of from about 5 to about 95 percent inventive composition.

Suitable binders include starch, gelatin, natural sugars, corn sweeteners, natural and synthetic gums such as acacia, sodium alginate, carboxymethylcellulose,
25 polyethylene glycol and waxes. Among the lubricants, there may be mentioned for use in these dosage forms, boric acid, sodium benzoate, sodium acetate, sodium chloride, and the like. Disintegrants include starch, methylcellulose, guar gum and the like. Sweetening and flavoring agents and preservatives may also be included where appropriate. Some of the terms noted above, namely
30 disintegrants, diluents, lubricants, binders and the like, are discussed in more detail below.

Additionally, the compositions of the present invention may be formulated in sustained release form to provide the rate controlled release of any one or more of
35 the components or active ingredients to optimize the therapeutic effects, i.e. antihistaminic activity and the like. Suitable dosage forms for sustained release include layered tablets containing layers of varying disintegration rates or controlled release polymeric matrices impregnated with the active components

and shaped in tablet form or capsules containing such impregnated or encapsulated porous polymeric matrices.

5 Liquid form preparations include solutions, suspensions and emulsions. As an example may be mentioned water or water-propylene glycol solutions for parenteral injections or addition of sweeteners and opacifiers for oral solutions, suspensions and emulsions. Liquid form preparations may also include solutions for intranasal administration.

10 Aerosol preparations suitable for inhalation may include solutions and solids in powder form, which may be in combination with a pharmaceutically acceptable carrier such as inert compressed gas, e.g. nitrogen.

15 For preparing suppositories, a low melting wax such as a mixture of fatty acid glycerides such as cocoa butter is first melted, and the active ingredient is dispersed homogeneously therein by stirring or similar mixing. The molten homogeneous mixture is then poured into convenient sized molds, allowed to cool and thereby solidifies.

20 Also included are solid form preparations which are intended to be converted, shortly before use, to liquid form preparations for either oral or parenteral administration. Such liquid forms include solutions, suspensions and emulsions.

25 The inventive pyridylpyrimidine compounds of the present invention may also be deliverable transdermally. The transdermal compositions may take the form of creams, lotions, aerosols and/or emulsions and can be included in a transdermal patch of the matrix or reservoir type as are conventional in the art for this purpose.

30 The term capsule refers to a special container or enclosure made of methyl cellulose, polyvinyl alcohols, or denatured gelatins or starch for holding or containing compositions comprising the active ingredients. Hard shell capsules are typically made of blends of relatively high gel strength bone and pork skin gelatins. The capsule itself may contain small amounts of dyes, opaquing agents, plasticizers and preservatives.

35

Tablet means compressed or molded solid dosage form containing the active ingredients with suitable diluents. The tablet can be prepared by compression of

mixtures or granulations obtained by wet granulation, dry granulation or by compaction well known to a person skilled in the art.

5 Oral gels refers to the active ingredients dispersed or solubilized in a hydrophilic semi-solid matrix.

Powders for constitution refers to powder blends containing the active ingredients and suitable diluents which can be suspended in water or juices.

10 Suitable diluents are substances that usually make up the major portion of the composition or dosage form. Suitable diluents include sugars such as lactose, sucrose, mannitol and sorbitol, starches derived from wheat, corn rice and potato, and celluloses such as microcrystalline cellulose. The amount of diluents in the
15 composition can range from about 5 to about 95% by weight of the total composition, preferably from about 25 to about 75%, more preferably from about 30 to about 60% by weight.

The term disintegrants refers to materials added to the composition to help it break apart (disintegrate) and release the medicaments. Suitable disintegrants
20 include starches, "cold water soluble" modified starches such as sodium carboxymethyl starch, natural and synthetic gums such as locust bean, karaya, guar, tragacanth and agar, cellulose derivatives such as methylcellulose and sodium carboxymethylcellulose, microcrystalline celluloses and cross-linked microcrystalline celluloses such as sodium croscarmellose, alginates such as
25 alginic acid and sodium alginate, clays such as bentonites, and effervescent mixtures. The amount of disintegrant in the composition can range from about 2 to about 20% by weight of the composition, more preferably from about 5 to about 10% by weight.

30 Binders characterize substances that bind or "glue" powders together and make them cohesive by forming granules, thus serving as the "adhesive" in the formulation. Binders add cohesive strength already available in the diluents or bulking agent. Suitable binders include sugars such as sucrose, starches derived from wheat, corn rice and potato; natural gums such as acacia, gelatin and
35 tragacanth; derivatives of seaweed such as alginic acid, sodium alginate and ammonium calcium alginate; cellulosic materials such as methylcellulose and sodium carboxymethylcellulose and hydroxypropylmethylcellulose; polyvinylpyrrolidone; and inorganics such as magnesium aluminum silicate. The

amount of binder in the composition can range from about 2 to about 20% by weight of the composition, more preferably from about 3 to about 10% by weight, even more preferably from about 3 to about 6% by weight.

- 5 Lubricant refers to a substance added to the dosage form to enable the tablet, granules, etc. after it has been compressed, to release from the mold or die by reducing friction or wear. Suitable lubricants include metallic stearates such as magnesium stearate, calcium stearate or potassium stearate; stearic acid; high melting point waxes; and water soluble lubricants such as sodium chloride,
10 sodium benzoate, sodium acetate, sodium oleate, polyethylene glycols and D,L-leucine. Lubricants are usually added at the very last step before compression, since they must be present on the surfaces of the granules and in between them and the parts of the tablet press. The amount of lubricant in the composition can range from about 0.2 to about 5% by weight of the composition, preferably from
15 about 0.5 to about 2%, more preferably from about 0.3 to about 1.5% by weight.

- Glidants are materials that prevent caking and improve the flow characteristics of granulations, so that flow is smooth and uniform. Suitable glidants include silicon dioxide and talc. The amount of glident in the composition can range from about
20 0.1% to about 5% by weight of the total composition, preferably from about 0.5 to about 2% by weight.

- Coloring agents are excipients that provide coloration to the composition or the dosage form. Such excipients can include food grade dyes and food grade dyes
25 adsorbed onto a suitable adsorbent such as clay or aluminum oxide. The amount of the coloring agent can vary from about 0.1 to about 5% by weight of the composition, preferably from about 0.1 to about 1%.

- As used herein, a "pharmaceutically effective amount" of an inhibitor and/or an
30 activator is an amount effective to achieve the desired physiological result, either in cells treated *in vitro* or in a subject treated *in vivo*. Specifically, a pharmaceutically effective amount is an amount sufficient to inhibit and or activate, for some period of time, one or more of the clinically defined pathological processes associated with the prion infection. The effective amount may vary
35 depending on the specific inhibitor and/or activator selected, and is also dependent on a variety of factors and conditions related to the subject to be treated and the severity of the infection. For example, if an inhibitor and/or activator is to be administered *in vivo*, factors such as the age, weight and health

of the patient as well as dose response curves and toxicity data obtained in pre-clinical animal work would be among those considered. If the inhibitor and/or activator is to be contacted with the cells *in vitro*, one would also design a variety of pre-clinical *in vitro* studies to assess such parameters as uptake, half-life, dose, toxicity, etc. The determination of a pharmaceutically effective amount for a given pharmaceutically active agent is well within the ability of those skilled in the art.

It is also apparent to a person skilled in the art that detection includes any method known in the art useful to indicate the presence, absence, or amount of a detection target. Such methods may include, but are not limited to, any molecular or cellular techniques, used singularly or in combination, including, but not limited to: hybridization and/or binding techniques, including blotting techniques and immunoassays; labeling techniques (chemiluminescent, colorimetric, fluorescent, radioisotopic); spectroscopic techniques; separations technology, including precipitations, electrophoresis, chromatography, centrifugation, ultrafiltration, cell sorting; and enzymatic manipulations (e.g., digestion).

It should be stressed that all above-mentioned features, aspects, and details of the present invention discussed and described in connection with infections and infectious diseases, equally apply to neurodegenerative diseases, like Alzheimer.

It is readily apparent to those skilled in the art that other suitable modifications and adaptations of the compositions and methods of the invention described herein are evident and may be made without departing from the scope of the invention or the embodiments disclosed herein. Having now described the present invention in detail, the same will be more clearly understood by reference to the following examples, which are included for purposes of illustration only and are not intended to be limiting of the invention.

30

Description of figures

Fig. 1 shows 6 selected pyridylpyrimidine derivatives which are suitable inhibitors for prion diseases, namely compounds 4, 5, 37, 52, 84, and 88;

Fig. 2 shows the compound 4-(4-Methylpiperazin-1-ylmethyl)-N-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide, also known as GleevecTM;

Fig. 3 shows selected compounds that have been identified as potent inhibitors in a prion propagation assay at a concentration of 5 μ m.

Examples

Materials and methods

5

1. Generation of cDNA-arrays on membranes

In order to manufacture cDNAs-arrays on membranes, the following strategy was pursued: cDNAs encoding parts of or full length proteins of interest – in the following referred to as “target cDNAs” – were cloned into the plasmid Bluescript II KS⁺ (Stratagene, USA). Large scale purifications of these plasmids were performed according to standard techniques and 200 µl aliquots (1 µg/µl plasmid concentration) were transferred into appropriate 96well plates. Plates were closed with sealing tape and chilled on ice for 5 minutes after incubation for 10 minutes at 95°C. 10 µl of 0.6 N NaOH were added and the mix was stored for 20 minutes at room temperature before addition of 10 µl 2.5 M Tris-HCl pH 7.1 and 20 µl 40x SSC (3 M NaCl, 300 mM Sodium Citrate, pH 7.0). Target cDNAs were spotted onto Nylon or Nitrocellulose membranes using a BioGrid (BioRobotics, UK) equipped with a 0.7 mm pintool. In this way, between 200 ng and 350 ng of plasmids encoding target cDNAs were transferred onto the membranes and crosslinked to the membranes by ultraviolet light (1.2×10^5 µJ/cm²). The arrays were stored for use in subsequent experiments at room temperature.

25

2. Generation of cells

PrP^{Sc}- and PrP^C-transfected mouse neuronal cells (N2A) were cultured in MEM (Minimum Essential Medium, Life Technologies) supplemented with 10% fetal calf serum at 37°C and 5% CO₂ to obtain $\sim 6 \times 10^6$ cells per tissue culture flask.

30

3. Lysis of cells, isolation of total RNA and purification of polyA⁺ RNA

After incubation of the cells with the virus for the respective time-points, cells were washed twice with phosphate buffered saline (PBS) and then trypsinized. Subsequently, cells were removed from the culture dish by resuspension with PBS. Afterwards, cells were sedimented and directly lysed in Tri reagent by repetitive pipetting using in 1ml of Tri reagent (Molecular Research Centre, Inc., USA) per 1×10^6 cells.

The lysates were stored at room temperature for 5 minutes and then centrifuged at 12000xg for 15 minutes at 4°C. The supernatant was mixed with 0,1 ml of 1-bromo-3-chloropropane per 1 ml of Tri reagent and vigorously shaken. The suspension was stored for 5 minutes at room temperature and then centrifuged at
5 12000xg for 15 minutes at 4°C.

The colourless upper phase was transferred into new tubes, mixed with 5 µl of poly-acryl-carrier (Molecular Research Centre, Inc., USA) and with 0.5 ml of isopropanol per 1 ml of Tri reagent and vigorously shaken. The samples were
10 stored at room temperature for 5 minutes and then centrifuged at 12000xg for 8 minutes at 4°C. The supernatant was removed and the RNA pellet washed twice with 1 ml of 75% ethanol. The pellet was dried and resuspended for 10 minutes at 55°C in 50 µl of RNase-free buffer (5 mM Tris-HCl pH 7.5). The integrity of the isolated RNA was determined by agarose/formaldehyde gel electrophoresis and
15 the RNA was finally stored at -70°C for use in subsequent experiments.

4. Preparation of radioactively labelled cDNA probes from RNA

In order to obtain radioactively labelled cDNA probes total RNA was transcribed
20 into a cDNA-probe in the presence of radioactively labelled dATP. 12 µl bidistilled DEPC (Diethylpyrocarbonate) treated H₂O containing 0.5 µg of primer TXN (5'-TTT TTT TTT TTT TTT TXN-3' with T → dTTP; N → dATP, dCTP, dGTP or dTTP; X → dATP, dCTP or dGTP) and total RNA (1 to 10 µg) were shaken between 5 and 15' at 60°C and then incubated on ice for 2 minutes. After
25 centrifugation (30 seconds, 10000xg) 7 µl of a mix consisting of 100 µCi dATP-P³³ (Amersham, UK) which were dried under vacuum previously and resuspended in 4 µl first strand buffer (Life Technologies, USA), 2 µl 0.1M DTT (Dithiothreitol) and 1 µl labelling solution (4 mM dCTP, dGTP, dTTP each and 80 µM dATP final concentration) were added. Following the addition of 1 µl Superscript II reverse
30 transcriptase (Life Technologies, USA) the reaction was incubated for 10 minutes at room temperature and then for 60 minutes at 38°C. Subsequently, the reaction was vigorously shaken for 30 minutes at 68°C after adding 5 µl 0.5 M EDTA and 25 µl 0.6M NaOH.

35 Unincorporated nucleotides were removed from the labelling reaction using ProbeQuant G-50 columns (Amersham, UK). The column was vigorously shaken and centrifuged for 1 minute at 735xg in an appropriate reaction tube after bottom closure and lid were removed. The column was placed into a new reaction tube,

the probe was applied onto the centre of the column material and the column was centrifuged for 2 minutes at 735xg. The flow-trough was transferred into new reaction tubes and filled up to a volume of 100 µl with bidistilled H₂O. The probe was precipitated by centrifugation for 15 minutes at 12000xg after 4 µl 5M NaCl, 1
5 µl poly-acryl-carrier (Molecular Research Centre, Inc., USA) and 250 µl ethanol were added. The supernatant was discarded and the pellet was dried at 50°C for 5 minutes before starting with the hybridisation.

5. Hybridisation of radioactively labelled cDNA-probes to cDNA-arrays

10

The pellet was resuspended in 10 µl C₀T DNA (1 µg/µl, Roche Diagnostics, Germany), 10 µl yeast tRNA (1µg/µl Sigma, USA) and 10 µl polyA (1 µg/µl, Roche Diagnostics, Germany) and incubated at 55°C for 5 minutes. Herring sperm DNA was added to a final concentration of 100 µg/ml and the volume was filled up to
15 100 µl with 5 µl 10% SDS (Sodiumdodecylsulfat), 25 µl 20x SSPE (3M Sodium chloride, 0,2 M Sodium dihydrogen phosphate monohydrate, 0,02 M Ethylenedinitrilo tetraacetic acid, disodium salt dihydrate; pH 7,4) and bidistilled H₂O. The mix was put on 95°C for 5 minutes, centrifuged for 30 seconds at 10000xg and vigorously shaken for 60 minutes at 65°C. A 1 µl aliquot of the
20 probe was used to measure the incorporation of radioactive dATP with a scintillation counter. Probes with at least a total of 20x10⁶ cpm were used.

The arrays were prehybridised for at least 3 hours at 42°C in hybridisation solution in a roller bottle oven. After prehybridization the radioactively labelled probe was
25 added into the hybridisation solution and hybridisation was continued for 20 to 40 hours.

The probe was discarded and replaced with wash solution A (2xSSC). The arrays were washed twice in wash solution A at room temperature in the roller
30 oven. Afterwards, wash solution A was replaced by wash solution B (2x SSC, 0.5% SDS) preheated to 65°C and arrays were washed twice for 30 minutes at 65°C. Then, wash solution B was replaced by wash solution C (0.5x SSC, 0.5% SDS) preheated to 65°C and arrays were washed twice for 30 minutes at 65°C. The moist arrays were wrapped in airtight bags and exposed for 8 to 72 hours on
35 erased phosphorimager screens (Fujifilm, Japan).

6. Analysis of cDNA-arrays

The exposed phosphorimager screens were scanned with a resolution of 100 μ and 16bits per pixel using a BAS-1800 (Fujifilm, Japan). Files were imported into the computer program ArrayVision (Imaging Research, Canada). Using the program's features, the hybridization signals of each target cDNA were converted into numbers. The strength of the hybridization signals reflected the quantity of RNA molecules present in the probe. Differentially expressed genes were selected according to the ratio of their signal strength after normalization to the overall intensity of the arrays.

7. Cell culture and expression of 3F4-tagged PrP (3F4-ScN2a)

The mouse neuroblastoma cell line 3F4-ScN2a represents a stably transfected clone of ScN2a cells (PrP^{Sc} infected N2a cells) which overexpress 3F4-epitope-tagged murine PrP. Residues 109 and 112 of murine PrP were replaced by methionine to introduce the epitope for reactivity with the monoclonal anti-PrP antibody 3F4. Cells were maintained in Dulbecco's modified Eagle's (DMEM) or Opti-MEM medium containing 10 % fetal calf serum, antibiotics and glutamin. For generation of stable transfectants we used the vector pcDNA3.1/Zeo (Invitrogen; Leek, The Netherlands). Lipofection of cells with recombinant plasmids was done using standard procedures and recombinant clones were selected by addition of 300 μ g Zeocin/ml medium.

8. Treatment of cells with inhibitors

All tested compounds were solubilized in DMSO (dimethylsulfoxide), and prepared as 10 mM stock solutions. The drugs were applied to the cells described above for three days in final concentrations between 5 and 20 μ M.

9. Immunoblot and proteinase K (PK) analysis

Confluent cell cultures were lysed in cold lysis buffer (10 mM Tris-HCl, pH 7.5; 100 mM NaCl; 10 mM EDTA; 0.5 % Triton X-100; 0.5 % DOC) (EDTA: ethylene diamine tetraacetate; Triton X-100: t-octylphenoxypolyethoxyethanol; DOC: deoxycholic acid). Postnuclear lysates were split between those with and without proteinase K digestion. Samples without proteinase K digestion were supplemented with proteinase inhibitors (5 mM PMSF, 0.5 mM Pefabloc, and aprotinin) (PMSF: phenylmethylsulfonyl fluoride) and directly precipitated with ethanol. Samples for proteinase K digestion were incubated with 20 μ g/ml proteinase K for 30 min at 37°C; digestion was stopped with proteinase inhibitors,

and samples were ethanol precipitated. After centrifuging for 30 min at 3,500 rpm the pellets were redissolved in TNE buffer (10 mM Tris-HCl pH7.5, 100 mM NaCl, 1mM EDTA) and gel loading buffer was then added. After boiling for 5 min an aliquot was analyzed on 12.5 % PAGE. For Western blot analysis, the proteins were electrotransferred to PVDF membranes (polyvinylidene difluoride). The membrane was blocked with 5 % non-fat dry milk in TBST (0.05 % Tween 20, 100 mM NaCl, 10 mM Tris-HCl, pH 7.8) (Tween 20: polyoxyethylenesorbitan monolaurate; Tris-HCl: Tris-(hydroxymethyl)-aminomethane-hydrochloride), incubated overnight with the primary antibody at 4°C and stained using the enhanced chemiluminescence blotting kit from Amersham Corporation. Specific immuno-staining of the PrP^C and PrP^{Sc} forms were obtained with the prion protein specific antibody 3F4 (Signet Pathologies, U.S.A.).

10. Results

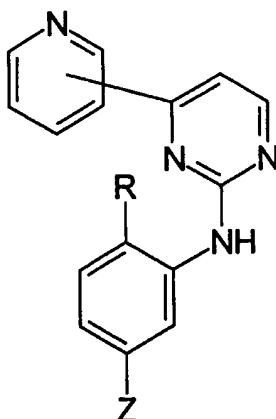
Determination of the amount of the pathogenic form of the prion protein PrP^{Sc} upon treatment of prion infected cells with different types of small molecule protein kinase inhibitors resulted in the identification of a compound class of pyridylpyrimidine derivatives exemplified by the compound 4-(4-Methylpiperazin-1-ylmethyl)-N-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide (compound 53) and compounds 4, 5, and 37.

These compounds significantly reduced the amount of PrP^{Sc} in prion infected cells in a concentration range between 5 and 20 µM (final concentration). As shown in Fig. 3 the selected compounds 4, 5, 37, and 53 inhibit almost completely the activity of prion propagation within said concentration range.

The compounds did not show any toxic effects on the cells in these concentrations. Therefore these molecules described herein serve as potential inhibitors for the medical intervention of prion diseases such as transmissible spongiform encephalitis (TSE) infections which include Bovine spongiform encephalitis (BSE) or the new variant of Creutzfeldt Jakob disease (vCJK).

Claims

1. Compounds having the general formula (I):



5

wherein:

R represents hydrogen or methyl;

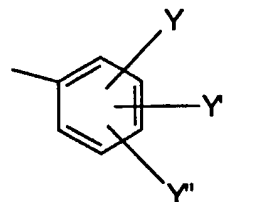
Y, Y', Y'' are independently of each other -H, -F, -Cl, -Br, -I, -CH₂F, -CH₂Cl, -CH₂Br, -CH₂I, -OH, -OCH₃, -CH₃, -CN, -OCF₃, 4-methylpiperazin-1-yl-methyl, -C(CH₃)=N-NH-C(NH)-NH₂;

10

Z represents -NO₂, -NH₂, -NH-CO-X, -NH-CS-X, -NH-CO-NH-X, -NH-SO₂-X;

X represents thiophenyl, cyclohexyl, isoquinolinyl, naphthyl, quinolinyl,

cyclopentyl, pyridinyl, naphthyridinyl, or

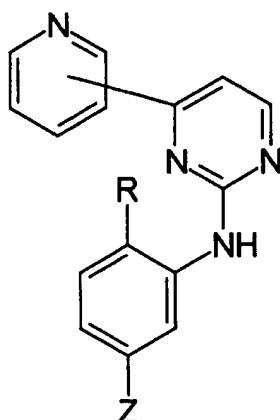


15

and pharmaceutically acceptable salts thereof.

20

2. Use of a compound having the general formula (I):



5

wherein:

R represents hydrogen or methyl;

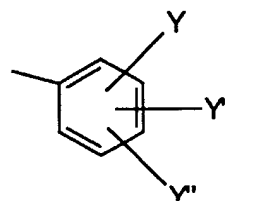
Y, Y', Y'' are independently of each other -H, -F, -Cl, -Br, -I, -CH₂F, -CH₂Cl, -CH₂Br, -CH₂I, -OH, -OCH₃, -CH₃, -CN, -OCF₃, 4-methylpiperazin-1-yl-methyl, -C(CH₃)=N-NH-C(NH)-NH₂;

10

Z represents -NO₂, -NH₂, -NH-CO-X, -NH-CS-X, -NH-CO-NH-X, -NH-SO₂-X;

X represents thiophenyl, cyclohexyl, isoquinolinyl, naphthyl, quinolinyl,

cyclopentyl, pyridinyl, naphthyridinyl, or

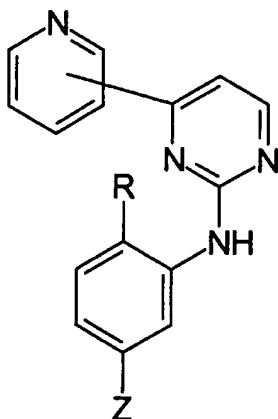


and pharmaceutically acceptable salts thereof as pharmaceutically active agents.

15

20

3. Use of a compound having the general formula (I):



5

wherein:

R represents hydrogen or methyl;

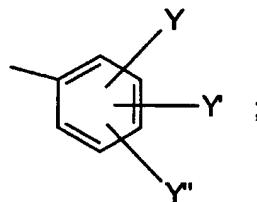
Y, Y', Y'' are independently of each other -H, -F, -Cl, -Br, -I, -CH₂F, -CH₂Cl, -CH₂Br, -CH₂I, -OH, -OCH₃, -CH₃, -CN, -OCF₃, 4-methylpiperazin-1-yl-methyl, -C(CH₃)=N-NH-C(NH)-NH₂;

10

Z represents -NO₂, -NH₂, -NH-CO-X, -NH-CS-X, -NH-CO-NH-X, -NH-SO₂-X;

X represents thiophenyl, cyclohexyl, isoquinolinyl, naphthyl, quinolinyl,

cyclopentyl, pyridinyl, naphthyridinyl, or



and pharmaceutically acceptable salts thereof for prophylaxis and/or treatment of infectious diseases or neurodegenerative diseases.

15

4. Use of a compound according to claim 2 or 3 for the prophylaxis and/or treatment of prion infections and/or diseases induced by prion infection.

20

5. Use of a compound according to any one of claims 2 - 4 wherein R represents hydrogen.
6. Use of a compound according to any one of claims 2 - 5 wherein Z represents -NH-CO-X or -NH-SO₂-X.

7. Use of a compound according to any one of claims 2 – 6 wherein Y, Y', Y'' are independently of each other –H, –F, –Cl, –CH₂F, –CH₂Cl, –OH, –OCH₃, –CH₃, –CN, –OCF₃, 4-methylpiperazin-1-yl-methyl.
- 5
8. Use of a compound according to claim 2 or 3 wherein the compound is selected from the group comprising:
- (3-Nitrophenyl)-(4-pyridin-3-yl-pyrimidin-2-yl)-amine;
- (3-Aminophenyl)-(4-pyridin-3-yl-pyrimidin-2-yl)-amine;
- 10 (5-Amino-2-methylphenyl)-(4-pyridin-3-yl-pyrimidin-2-yl)-amine;
- 4-Chloromethyl-N-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 4-Chloromethyl-N-[3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 4-(4-Methylpiperazin-1-ylmethyl)-N-[3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 15 Thiophene-3-carboxylic acid [4-methyl-3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- 4-Chloro-N-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 20 4-Chloro-N-[3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 3,4,5-Trimethoxy-N-[3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 4-Cyano-N-[3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 4-Methoxy-N-[3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 25 4-Chloro-N-[3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzenesulfonamide;
- Thiophene-3-carboxylic acid [3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- 3,5-Dimethoxy-N-[3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 30 3,4,5-Trimethoxy-N-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 4-Cyano-N-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 4-Methoxy-N-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 35 4-Chloro-N-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzenesulfonamide;

- Thiophene-3-carboxylic acid [4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- 3,5-Dimethoxy-*N*-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 5 4-Trifluoromethoxy-*N*-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- Cyclohexanecarboxylic acid [4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- 10 Cyclohexanecarboxylic acid [3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- Isoquinoline-5-sulfonic acid [4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- Isoquinoline-5-sulfonic acid [3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- 15 (5-Nitro-2-methylphenyl)-(4-pyridin-2-yl-pyrimidin-2-yl)-amine;
- (5-Amino-2-methylphenyl)-(4-pyridin-2-yl-pyrimidin-2-yl)-amine;
- 3,4,5-Trimethoxy-*N*-[4-methyl-3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 4-Cyano-*N*-[4-methyl-3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 20 (3-Aminophenyl)-(4-pyridin-2-yl-pyrimidin-2-yl)-amine;
- 4-Chloro-*N*-[4-methyl-3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- Cyclohexanecarboxylic acid [4-methyl-3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- 25 4-Cyano-*N*-[4-methyl-3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 4-Chloro-*N*-[4-methyl-3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzenesulfonamide;
- 30 4-Methoxy-*N*-[4-methyl-3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 4-Chloro-*N*-[4-methyl-3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- Cyclohexanecarboxylic acid [3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- 35 3,5-Dimethoxy-*N*-[3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- (5-Amino-2-methylphenyl)-(4-pyridin-4-yl-pyrimidin-2-yl)-amine;

- Thiophene-3-carboxylic acid [3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- 4-Chloro-*N*-[3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzenesulfonamide;
- 5 4-Chloro-*N*-[3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- (3-Aminophenyl)-(4-pyridin-4-yl-pyrimidin-2-yl)-amine;
- (3-Nitrophenyl)-(4-pyridin-4-yl-pyrimidin-2-yl)-amine;
- 4-Trifluoromethoxy-*N*-[4-methyl-3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 10 Isoquinoline-5-sulfonic acid [3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- 4-Methoxy-*N*-[3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 4-Cyano-*N*-[3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 3,4,5-Trimethoxy-*N*-[3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 15 3,5-Dimethoxy-*N*-[4-methyl-3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 3,4,5-Trimethoxy-*N*-[4-methyl-3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 20 4-(4-Methylpiperazin-1-ylmethyl)-*N*-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 4-Methyl-*N*-[4-methyl-3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzenesulfonamide;
- 4-Methoxy-*N*-[4-methyl-3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 25 3,5-Dimethoxy-*N*-[4-methyl-3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- Naphthalene-2-carboxylic acid [4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- 30 *N*-[3-(4-Pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 4-Chloro-*N*-[3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 4-Methoxy-*N*-[3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 4-Chloro-*N*-[3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzenesulfonamide;
- 35 Thiophene-2-carboxylic acid 3-(4-pyridin-2-yl-pyrimidin-2-yl-amino)-phenyl]-amide;
- Naphthalene-2-sulfonic-acid [3-(4-pyridin-2-yl-pyrimidin-2-yl-amino)-phenyl]-amide;

- Isoquinoline-5-sulfonic-acid [3-(4-pyridin-2-yl-pyrimidin-2-yl-amino)-phenyl]-amide;
- Cyclopentanecarboxylic acid 3-(4-pyridin-2-yl-pyrimidin-2-yl-amino)-phenyl]-amide;
- 5 Naphthalene-2-carboxylic acid [3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- 4-Cyano-*N*-[3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 3,5-Dimethoxy-*N*-[3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 4-Bromo-*N*-[3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 10 4-Methyl-*N*-[3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 4-Fluoro-*N*-3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzenesulfonamide;
- 3,5-Dichloro-*N*-[3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- N*-[3-(4-Pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 15 4-Chloromethyl-*N*-[3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 4-Methyl-*N*-3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzenesulfonamide;
- 4-(4-Methylpiperazin-1-ylmethyl)-*N*-[3-(4-pyridin-2-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 20 Naphthalene-2-carboxylic acid [3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- 2-Methoxy-*N*-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 2-Methoxy-*N*-[3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 25 4-Methyl-*N*-[3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 4-Methyl-*N*-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- N*-[4-Methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
- 1-(3,5-Diacetyl-phenyl)-3-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-urea;
- 30 1-{3,5-Bis-(amidinohydrazone)-phenyl}-3-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-urea;
- N*-[4-Methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-nicotinamide;
- N*-[3-(4-Pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-nicotinamide;
- 35 [1,8]Naphthyridine-2-carboxylic acid [3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-amide;
- [1,8]Naphthyridine-2-carbothioic acid [3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-amide;

- 2-Methoxy-*N*-[3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
4-Trifluoromethoxy-*N*-[3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-
benzamide;
4-Methyl-*N*-[3-(4-pyridin-4-yl-pyrimidin-2-ylamino)-phenyl]-benzamide;
5 and/or a pharmaceutically acceptable salt of these compounds.
9. Use according to claim 8 wherein the compound is 4-(4-Methylpiperazin-1-ylmethyl)-*N*-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-ylamino)-phenyl]-benzamide.
10. Use of a compound recited in any one of claims 2 – 9 and/or pharmaceutically acceptable salts thereof for the manufacture of a pharmaceutical composition for prophylaxis and/or treatment of prion infections and/or diseases induced by prion infection and/or neurodegenerative diseases.
11. Use according to claim 4 or 10 wherein said prion infection and/or disease is selected from the group comprising Scrapie, TME, CWD, BSE, CJD, vCJD, GSS, FFI, Kuru, and Alpers Syndrome.
12. Use according to claim 11 wherein said prion infection is BSE, vCJD, or CJD.
13. Use of a compound recited in any one of claims 2 – 9 as an inhibitor for at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, GPIR-1.
14. Use of a compound according to any one of claims 2 to 13 wherein the compound of the general formula (I) and/or pharmaceutically acceptable salts thereof is administered in a dosage corresponding to an effective concentration in the range of 0.01 – 50 μ M.
15. Pharmaceutical composition comprising at least one compound recited in any one of claims 2 – 9 as an active ingredient, together with one or more pharmaceutically acceptable carrier(s), excipient(s) or diluents.

16. Method for preventing and/or treating infections and/or diseases in an individual which comprises administering to the individual an amount of at least one compound recited in claims 2 – 9 and/or pharmaceutically acceptable salts thereof effective to prevent and/or treat said infections and/or diseases.
17. Method for preventing and/or treating prion infections and/or prion diseases induced by prion infections in an individual which comprises administering to the individual an amount of at least one compound recited in any one of claims 3 to 8 and/or pharmaceutically acceptable salts thereof effective to prevent and/or treat said prion infection and/or disease.
18. Method for preventing and/or treating prion infections and/or prion diseases induced by prion infections in an individual which comprises administering to the individual an amount of at least one compound recited in claim 8 and/or pharmaceutically acceptable salts thereof effective to prevent and/or treat said prion infection and/or disease.
19. Method for detecting prion infections and/or prion diseases in an individual comprising:
- a) providing a sample from said individual;
 - b) adding to said sample a pharmaceutically effective amount of at least one pharmaceutically active agent; and
 - c) detecting activity in said sample of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, GPIR-1.
20. Method according to claim 19 wherein said sample comprises blood, milk, saliva, sputum, excrement, urine, spinal cord liquid, liquor, lachrymal gland liquid, biopsies and all other samples that can be taken from a living animal or human for diagnostic purposes.
21. Method for detecting prion infections and/or prion diseases in cells, cell cultures and/or cell lysates comprising:
- a) providing said cells, cell cultures and/or cell lysates;
 - b) adding to said cells, cell cultures and/or cell lysates a pharmaceutically effective amount of at least one pharmaceutically active agent; and

- c) detecting activity in said sample of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, GPIR-1.

5

22. Method for preventing and/or treating prion infections and/or prion diseases in an individual comprising the step of administering a pharmaceutically effective amount of at least one pharmaceutically active agent which inhibits at least partially the activity of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, GPIR-1, or which inhibits at least partially the production of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, GPIR-1.

10

15

23. Method for preventing and/or treating prion infections and/or prion diseases in cell or cell cultures comprising the step of administering a pharmaceutically effective amount of at least one pharmaceutically active agent which inhibits at least partially the activity of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, GPIR-1, or which inhibits at least partially the production of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, GPIR-1.

20

25

24. Method for regulating the production of prions in an individual comprising the step of administering a pharmaceutically effective amount of at least one pharmaceutically active agent which inhibits at least partially the activity of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, GPIR-1, or which inhibits at least partially the production of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt,

30

35

Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, GPIR-1.

- 5 25. Method for regulating the production of prions in cells comprising the step of administering a pharmaceutically effective amount of at least one pharmaceutically active agent which inhibits at least partially the activity of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, 10 HSP86, GPIR-1, or which inhibits at least partially the production of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, GPIR-1.
- 15 26. A monoclonal or polyclonal antibody that binds to a human cellular protein kinase, phosphatase or a cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1.
- 20 27. Method according to any one of claims 19 – 25, wherein the agent is a monoclonal or polyclonal antibody which binds to a human cellular protein kinase, phosphatase or a cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, 25 JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1.
- 30 28. Method according to any one of claims 19 – 25, wherein the agent is at least one compound of the general formula (I) and/or pharmaceutically acceptable salts thereof.
- 35 29. Method according to any one of claims 16 – 25, wherein the agent is 4-(4-Methylpiperazin-1-ylmethyl)-N-[4-methyl-3-(4-pyridin-3-yl-pyrimidin-2-yl-amino)-phenyl]-benzamide and/or pharmaceutically acceptable salts thereof.
- 30 30. Method according to claim 28 wherein the compound of the general formula (I) and/or pharmaceutically acceptable salts thereof is administered in a dosage corresponding to an effective concentration in the range of 0.01 – 50 μ M.

31. Method for detecting compounds useful for the prophylaxis and/or treatment of prion infections and/or diseases comprising:
- 5 a) contacting a test compound with at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, GPIR-1; and
- 10 b) detecting the activity of said human cellular protein kinase, phosphatase or cellular signal transduction molecule.
32. Method for preventing and/or treating prion infections and/or diseases in an individual comprising the step of administering a pharmaceutically effective amount of at least one pharmaceutically active agent which activates at least partially the activity of at least one human cellular protein kinase,
- 15 phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1, or which activates or stimulates the production of at least one human cellular protein kinase,
- 20 phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1.
33. Method for regulating the production of prions in an individual comprising the step of administering an individual a pharmaceutically effective amount of at least one pharmaceutically active agent wherein said agent activates at least partially the activity of at least one human cellular protein kinase,
- 25 phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1, or wherein said agent at least partially activates or stimulates the production of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1.
- 30
- 35
34. Method for regulating the production of prions in cells comprising the step of administering the cells a pharmaceutically effective amount of at least one pharmaceutically active agent wherein said agent activates at least partially

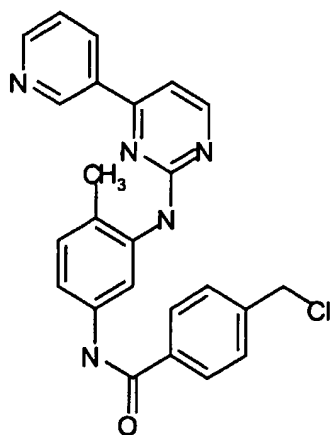
- the activity of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1, or wherein said agent at least partially
- 5 activates or stimulates the production of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1 in the cells.
- 10 35. Method for regulating the expression of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1 in an
- 15 individual comprising the step of administering the individual a pharmaceutically effective amount of at least one pharmaceutically active agent wherein said agent inhibits at least partially the transcription of DNA or the translation of RNA.
- 20 36. Method for regulating the expression of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1 in the cells comprising the step of administering the cells a pharmaceutically effective
- 25 amount of at least one pharmaceutically active agent wherein said agent inhibits at least partially the transcription of DNA or the translation of RNA.
- 30 37. Oligonucleotide that binds to the DNA or RNA encoding a human cellular protein kinase, phosphatase or a cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1.
- 35 38. Method according to claim 22, 23, 24, 25, 35 or 36 wherein the agent is a oligonucleotide which binds to the DNA and/or RNA encoding a human cellular protein kinase, phosphatase or a cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1.

39. Method according to claims 16, 17, 18, 19, 22, 24, 32, 33, or 35 wherein said individual is a human or ruminant.
- 5 40. Method according to any one of claims 17, 18, 19, 21, 22, 23, 31, or 32 wherein said prion infection and/or prion disease is selected from the group comprising Scrapie, TME, CWD, BSE, vCJD, CJD, GSS, FFI, Kuru, and Alpers Syndrome.
- 10 41. Method according to claim 40 wherein said prion infection and/or prion disease is BSE, vCJD, or CJD.
- 15 42. A solid support useful for detecting prion infections and/or diseases in an individual, the solid support comprising an immobilized oligonucleotide, wherein said oligonucleotide is capable of detecting activity of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1.
- 20 43. A solid support useful for detecting prion infections and/or diseases in cells, the solid support comprising an immobilized oligonucleotide, wherein said oligonucleotide is capable of detecting activity of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1.
- 25 44. A solid support useful for screening compounds useful for the prophylaxis and/or treatment of prion infections and/or diseases in an individual, the solid support comprising at least one immobilized oligonucleotide, wherein said oligonucleotide encodes one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1.
- 30 45. A solid support useful for screening compounds useful for the prophylaxis and/or treatment of prion infections and/or diseases in an individual, the solid support comprising at least one immobilized human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the

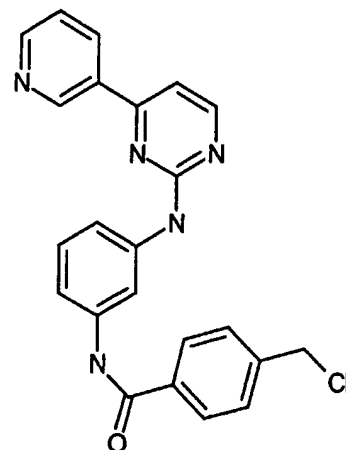
group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1.

- 5 46. Composition useful for the prophylaxis and/or treatment of an individual afflicted with prions comprising at least one agent capable of inhibiting at least partially the activity of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1.
- 10 47. Composition useful for the prophylaxis and/or treatment of an individual afflicted with prions comprising at least one agent capable of activating or stimulating at least partially the activity of at least one human cellular protein kinase, phosphatase or cellular signal transduction molecule selected from the group comprising FGF-R1, Tkt, Abl, clk1, MKK7, LIMK-2, CaM-KI, JNK2, CDC2, PRK, PTP-SL, PTP-zeta, HSP86, and GPIR-1.
- 15 48. Composition according claim 46 or 47, wherein the agent is at least one compound of the general formula (I) and/or pharmaceutically acceptable salts thereof.
- 20

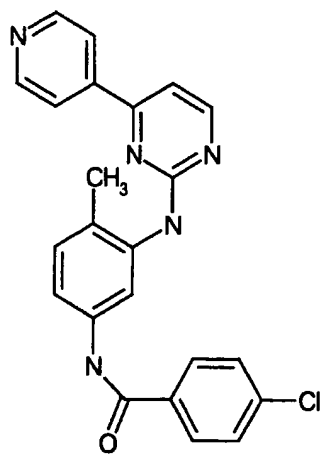
Fig. 1



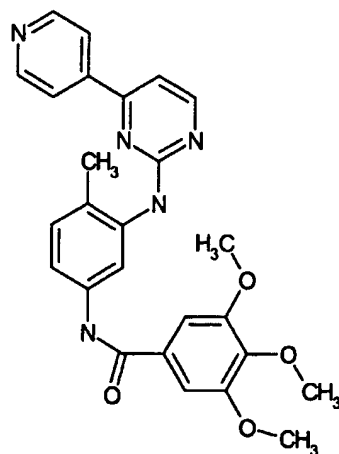
Compound 4



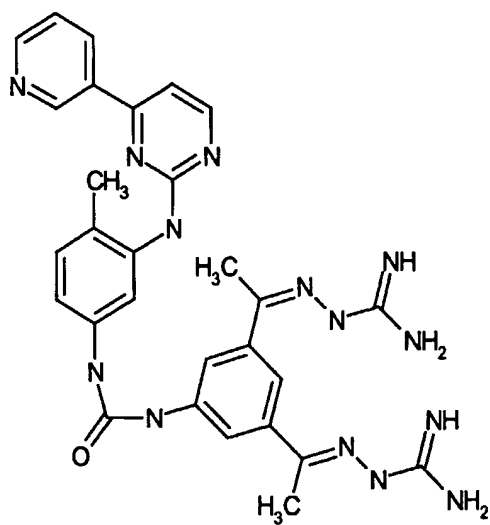
Compound 5



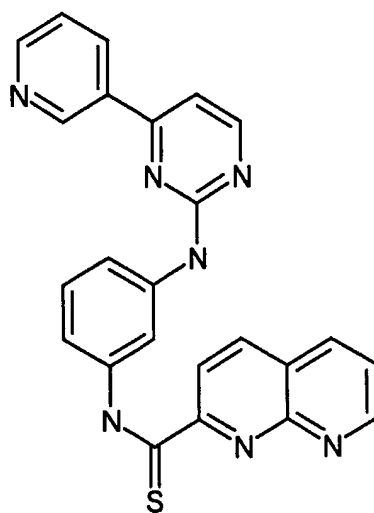
Compound 37



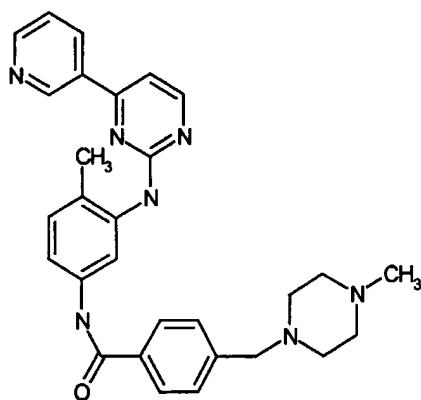
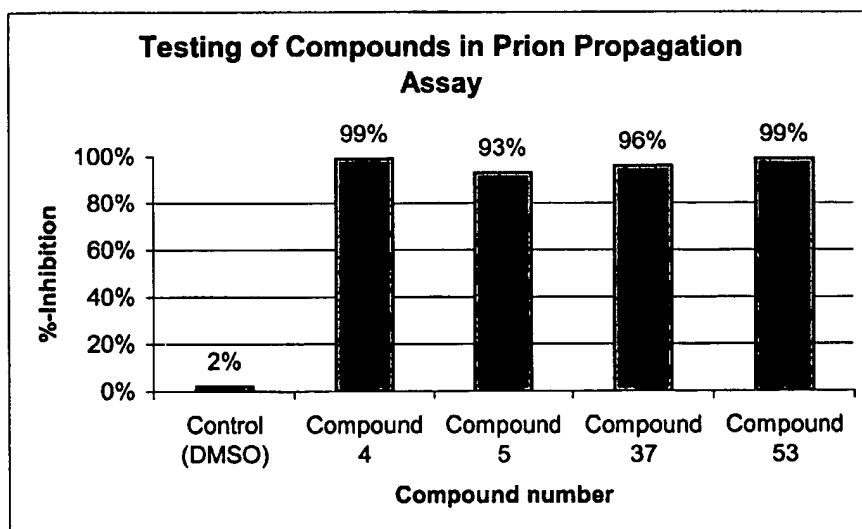
Compound 52



Compound 84



Compound 88

Fig. 2**Compound 53 (Gleevec™)****Fig. 3**

SEQUENCE LISTING

<110> Axxima Pharmaceuticals AG

5 <120> Human cellular protein kinases and phosphatases as
targets for diagnosis and treatment of prion diseases

<130> AXX-P11001-WO

10 <140> 0111858.5
<141> 2001-05-16

<150> US 60/293,528
<151> 2001-05-29

15 <160>

<170> PatentIn Ver. 2.1

20 <210> 1
<211> 2662
<212> DNA
<213> Homo sapiens

25 <220> FGF-R1
<223> Description of Sequence: N/A

<400> 1

tcagtttgaa aaggaggatc gagctcactc gtggagtatc catggagatg tggagccttg 60

30 tcaccaacct ctaactgcag aactgggatg tggagctgga agtgcctcct cttctgggct 120

gtgctgggtca cagccacact ctgcaccgct aggccgtccc cgaccttgcc tgaacaagcc 180

cagccctggg gagcccctgt ggaagtggag tccttcctgg tccaccccggtgacctgctg 240

cagcttcgct gtcggctgctg ggacgatgtg cagagcatca actggctgctg ggacgggggtg 300

cagctggcgg aaagcaaccg caccgcctac acaggggagg aggtggagggt gcaggactcc 360

35 gtgccgcag actccggcct ctatgcttgc gtaaccagca gcccctcggg cagtgcacacc 420

acctacttct ccgtcaatgt ttcagatgct ctccctcct cggaggatga tgatgatgat 480

gatgactcct cttcagagga gaaagaaaca gataacacca aaccaaaccg tatgcccgtg 540

gtcccatatt ggacatcccc agaaaagatg gaaaagaaat tgcattgcagt gccggctgcc 600

aagacagtga agttcaaag cccttccagt gggaccccaa accccacact gcgctggttg 660
aaaaatggca aagaattcaa acctgaccac agaattggag gctacaaggt ccgttatgcc 720
acctggagca tcataatgga ctctgtggtg ccctctgaca agggcaacta cacctgcatt 780
gtggagaatg agtacggcag catcaaccac acataccagc tggatgtcgt ggagcggtcc 840
5 cctcaccgcc ccattcctgca agcagggttg cccgccaca aaacagtggc cctgggtagc 900
aacgtggagt tcatgtgtaa ggtgtacagt gaccgcagc cgcacatcca gtggctaaag 960
cacatcgagg tgaatgggag caagattggc ccagacaacc tgcttatgt ccagatcttg 1020
aagactgctg gagttaatac caccgacaaa gagatggagg tgcttcactt aagaaatgtc 1080
tcctttgagg acgcagggga gtatacgtgc ttggcgggta actctatcgg actctcccat 1140
10 cactctgcat ggttgaccgt tctggaagcc ctggaagaga ggccggcagt gatgacctcg 1200
cccctgtacc tggagatcat catctattgc acaggggcct tcctcatctc ctgcatggtg 1260
gggtcggtca tcgtctacaa gatgaagagt ggtaccaaga agagtgactt ccacagccag 1320
atggctgtgc acaagctggc caagagcatc cctctgcgca gacaggtaac agtgtctgct 1380
gactccagtg catccatgaa ctctggggtt cttctggttc ggccatcacg gctctcctcc 1440
15 agtgggactc ccattgctagc aggggtctct gagtatgagc ttcccgaaga ccctcgctgg 1500
gagctgcctc gggacagact ggtcttaggc aaaccctgg gagagggctg ctttgggagc 1560
gtggtgttgg cagaggttat cgggctggac aaggacaaac ccaaccgtgt gaccaaagtg 1620
gctgtgaaga tgttgaagtc ggacgcaaca gagaaagact tgtcagacct gatctcagaa 1680
atggagatga tgaagatgat cgggaagcat aagaatatca tcaacctgct gggggcctgc 1740
20 acgcaggatg gtcccttgta tgtcatcgtg gagtatgcct ccaagggcaa cctgcgggag 1800
tacctgcagg cccggaggcc cccagggtg gaatactgct acaaccacag ccacaacca 1860
gaggagcagc tctcctcaa ggacctggtg tctgcgcct accagggtggc ccgaggcatg 1920
gagtatctgg cctccaagaa gtgcatacac cgagacctgg cagccaggaa tgtcctggtg 1980
acagaggaca atgtgatgaa gatagcagac tttggcctcg cacgggacat tcaccacatc 2040
25 gactactata aaaagacaac caacggccga ctgcctgtga agtggatggc acccgaggca 2100
ttatttgacc ggatctacac ccaccagagt gatgtgtggt ctttcggggt gctcctgtgg 2160
gagatcttca ctctgggcgg ctcccatcac cccggtgtgc ctgtggagga acttttcaag 2220
ctgctgaagg aggtcaccg catggacaag cccagtaact gcaccaacga gctgtacatg 2280
atgatgcggg actgctggca tgcagtgcct tcacagagac ccacctcaa gcagctggtg 2340
30 gaagacctgg accgcatcgt ggccttgacc tccaaccagg agtacctgga cctgtccatg 2400
cccctggacc agtactcccc cagctttccc gacaccgga gctctacgtg ctctcaggg 2460
gaggattccg tcttctctca tgagccgctg cccgaggagc cctgcctgcc ccgacacca 2520
gccagcttg ccaatggcgg actcaaacgc cgctgactgc caccacacg ccctccccag 2580
actccaccgt cagctgtaac cctcaccac agccctgcc tggggccacc acctgtccgt 2640
35 ccctgtcccc tttcctgctg gg 2662

<211> 822

<212> PRT

<213> Homo sapiens

5 <220>

<223> Description of Sequence: N/A

<400> 2

Met Trp Ser Trp Lys Cys Leu Leu Phe Trp Ala Val Leu Val Thr Ala
 10 1 5 10 15
 Thr Leu Cys Thr Ala Arg Pro Ser Pro Thr Leu Pro Glu Gln Ala Gln
 20 25 30
 Pro Trp Gly Ala Pro Val Glu Val Glu Ser Phe Leu Val His Pro Gly
 35 40 45
 15 Asp Leu Leu Gln Leu Arg Cys Arg Leu Arg Asp Asp Val Gln Ser Ile
 50 55 60
 Asn Trp Leu Arg Asp Gly Val Gln Leu Ala Glu Ser Asn Arg Thr Arg
 65 70 75 80
 Ile Thr Gly Glu Glu Val Glu Val Gln Asp Ser Val Pro Ala Asp Ser
 20 85 90 95
 Gly Leu Tyr Ala Cys Val Thr Ser Ser Pro Ser Gly Ser Asp Thr Thr
 100 105 110
 Tyr Phe Ser Val Asn Val Ser Asp Ala Leu Pro Ser Ser Glu Asp Asp
 115 120 125
 25 Asp Asp Asp Asp Asp Ser Ser Ser Glu Glu Lys Glu Thr Asp Asn Thr
 130 135 140
 Lys Pro Asn Arg Met Pro Val Ala Pro Tyr Trp Thr Ser Pro Glu Lys
 145 150 155 160
 Met Glu Lys Lys Leu His Ala Val Pro Ala Ala Lys Thr Val Lys Phe
 30 165 170 175
 Lys Cys Pro Ser Ser Gly Thr Pro Asn Pro Thr Leu Arg Trp Leu Lys
 180 185 190
 Asn Gly Lys Glu Phe Lys Pro Asp His Arg Ile Gly Gly Tyr Lys Val
 195 200 205
 35 Arg Tyr Ala Thr Trp Ser Ile Ile Met Asp Ser Val Val Pro Ser Asp
 210 215 220
 Lys Gly Asn Tyr Thr Cys Ile Val Glu Asn Glu Tyr Gly Ser Ile Asn
 225 230 235 240

His Thr Tyr Gln Leu Asp Val Val Glu Arg Ser Pro His Arg Pro Ile
 245 250 255
 Leu Gln Ala Gly Leu Pro Ala Asn Lys Thr Val Ala Leu Gly Ser Asn
 260 265 270
 5 Val Glu Phe Met Cys Lys Val Tyr Ser Asp Pro Gln Pro His Ile Gln
 275 280 285
 Trp Leu Lys His Ile Glu Val Asn Gly Ser Lys Ile Gly Pro Asp Asn
 290 295 300
 Leu Pro Tyr Val Gln Ile Leu Lys Thr Ala Gly Val Asn Thr Thr Asp
 10 305 310 315 320
 Lys Glu Met Glu Val Leu His Leu Arg Asn Val Ser Phe Glu Asp Ala
 325 330 335
 Gly Glu Tyr Thr Cys Leu Ala Gly Asn Ser Ile Gly Leu Ser His His
 340 345 350
 15 Ser Ala Trp Leu Thr Val Leu Glu Ala Leu Glu Glu Arg Pro Ala Val
 355 360 365
 Met Thr Ser Pro Leu Tyr Leu Glu Ile Ile Ile Tyr Cys Thr Gly Ala
 370 375 380
 Phe Leu Ile Ser Cys Met Val Gly Ser Val Ile Val Tyr Lys Met Lys
 20 385 390 395 400
 Ser Gly Thr Lys Lys Ser Asp Phe His Ser Gln Met Ala Val His Lys
 405 410 415
 Leu Ala Lys Ser Ile Pro Leu Arg Arg Gln Val Thr Val Ser Ala Asp
 420 425 430
 25 Ser Ser Ala Ser Met Asn Ser Gly Val Leu Leu Val Arg Pro Ser Arg
 435 440 445
 Leu Ser Ser Ser Gly Thr Pro Met Leu Ala Gly Val Ser Glu Tyr Glu
 450 455 460
 Leu Pro Glu Asp Pro Arg Trp Glu Leu Pro Arg Asp Arg Leu Val Leu
 30 465 470 475 480
 Gly Lys Pro Leu Gly Glu Gly Cys Phe Gly Gln Val Val Leu Ala Glu
 485 490 495
 Ala Ile Gly Leu Asp Lys Asp Lys Pro Asn Arg Val Thr Lys Val Ala
 500 505 510
 35 Val Lys Met Leu Lys Ser Asp Ala Thr Glu Lys Asp Leu Ser Asp Leu
 515 520 525
 Ile Ser Glu Met Glu Met Met Lys Met Ile Gly Lys His Lys Asn Ile
 530 535 540

[illegible]

<210> 3

<211> 3840

<212> DNA

5 <213> Homo sapiens

<220> Abl

<223> Description of Sequence: N/A

10 <400> 3

ggccttcccc ctgcgaggat cgccgttggc ccgggttggc tttggaaagc ggcggtggct 60
ttggggccggg ctcgccctcg ggaacgccag gggcccctgg gtgcggacgg gcgcggccag 120
gaggggggta aggcgcaggc ggcggcgggg cggggcgggg cctggcgggc gccctctccg 180
ggccctttgt taacaggcgc gtcccggcca gcggagacgc ggccgccttg ggcgggcgcg 240
15 ggcggcgggc ggcggtgagg gcggcctgcg gggcggcgcc cgggggcccg gccgagccgg 300
gcctgagccg ggcccggacc gagctgggag aggggctccg gcccgatcgt tcgcttggcg 360
caaaaatgtg gagatctgcc tgaagctggt gggctgcaaa tccaagaagg ggctgtcctc 420
gtcctccagc tgttatctg aagaagccct tcagcggcca gtagcatctg actttgagcc 480
tcagggtctg agtgaagccg ctcgttggaa ctccaaggaa aaccttctcg ctggacceag 540
20 tgaaaatgac cccaaccttt tcgttgact gtatgatttt gtggccagtg gagataaac 600
tctaagcata actaaagggt aaaagctccg ggtcttaggc tataatcaca atggggaatg 660
gtgtgaagcc caaaccaaaa atggccaagg ctgggtccca agcaactaca tcacgccagt 720
caacagtctg gagaaacact cctggtacca tgggcctgtg tcccgaatg ccgctgagta 780
tccgctgagc agcgggatca atggcagctt cttggtgctg gagagtgaga gcagtcctag 840
25 ccagaggtcc atctcgctga gatacgaagg gaggtgtac cattacagga tcaacactgc 900
ttctgatggc aagctctacg tctctccga gagccgcttc aacaccctgg ccgagttggt 960
tcatcatcat tcaacggtgg ccgacgggct catcaccacg ctccattatc cagcccaaaa 1020
gcgcaacaag cccactgtct atggtgtgtc cccaactac gacaagtggg agatggaacg 1080
cacggacatc accatgaagc acaagctggg cgggggcccag tacggggagg tgtacgaggg 1140
30 cgtgtggaag aaatacagcc tgacggtggc cgtgaagacc ttgaaggagg acacatgga 1200
ggtggaagag ttcttgaaag aagctgcagt catgaaagag atcaaacacc ctaacctagt 1260
gcagctcctt ggggtctgca cccgggagcc cccgttctat atcatcactg agttcatgac 1320
ctacgggaac ctctggact acctgaggga gtgcaaccgg caggaggtga acgccgtggt 1380
gctgctgtac atggccactc agatctcgtc agccatggag tacctagaga agaaaaactt 1440
35 catccacaga gatcttgctg cccgaaactg cctggtaggg gagaaccact tgggtgaagg 1500
agctgatttt ggctgagca ggttgatgac aggggacacc tacacagccc atgctggagc 1560
caagttcccc atcaaagga ctgcaccgga gagcctggcc tacaacaagt tctccatcaa 1620
gtccgacgtc tgggcatttg gagtattgct ttgggaaatt gctacctatg gcatgtcccc 1680

ttacccgga attgaccgtt cccaggtgta tgagctgcta gagaaggact accgcatgaa 1740
gcgcccagaa ggctgcccag agaaggtcta tgaactcatg cgagcatgtt ggcagtggaa 1800
tccctctgac cggccctcct ttgctgaaat ccaccaagcc tttgaaacaa tgttccagga 1860
atccagtatc tcagacgaag tggaaaagga gctggggaaa caaggcgtcc gtggggctgt 1920
5 gactaccttg ctgcaggccc cagagctgcc caccaagacg aggacctcca ggagagctgc 1980
agagcacaga gacaccactg acgtgcctga gatgcctcac tccaagggcc agggagagag 2040
cgatcctctg gaccatgagc ctgcccgtgc tccattgctc cctcgaaaaag agcgagggtcc 2100
cccgaggggc ggctggaatg aagatgagcg ccttctcccc aaagacaaaa agaccaactt 2160
gttcagcgcc ttgatcaaga agaagaagaa gacagcccca acccctccca aacgcagcag 2220
10 ctcttcccg gagatggacg gccagccgga gcgcagaggg gccggcgagg aagagggccg 2280
agacatcagc aacggggcac tggctttcac cccttggac acagctgacc cagccaagtc 2340
cccaaagccc agcaatgggg ctggggctcc caatggagcc ctccgggagt cggggggtc 2400
aggcttcccg tctccccacc tgtggaagaa gtccagcacg ctgaccagca gccgcctagc 2460
caccggcgag gaggagggcg gtggcagctc cagcaagcgc ttctgctcgt cttgctccgt 2520
15 ctctgctgtt ccccatgggg ccaaggacac ggagtggagg tcagtcacgc tgctcggga 2580
cttgagctcc acgggaagac agtttgactc gtccacattt ggagggcaca aaagtgagaa 2640
gccggtctg cctcggaaga gggcagggga gaacaggtct gaccaggtga cccgaggcac 2700
agtaacgcct cccccaggc tgggaaaaa gaatgaggaa gctgctgatg aggtcttcaa 2760
agacatcatg gagtccagcc cgggtccag ccgcccac ctgactcaa aaccctccg 2820
20 gcggcaggtc accgtggccc ctgcctcggg cctccccac aaggaagaag cctggaaagg 2880
cagtgcctta gggaccctg ctgcagctga gccagtgacc cccaccagca aagcaggctc 2940
aggtgcacca aggggcacca gcaagggccc cgcgaggag tccagagtga ggaggcacia 3000
gcactcctct gagtccagc ggagggacaa ggggaaattg tccaagctca aacctgccc 3060
gccgccccca ccagcagcct ctgcagggaa ggctggagga aagccctcgc agaggcccg 3120
25 ccaggaggct gccggggagg cagtcttggg cgcaagaca aaagccacga gtctggttga 3180
tgctgtgaac agtgacgctg ccaagcccag ccagccggca gagggcctca aaaagcccg 3240
gtctccggcc actccaaagc cacacccgc caagccgtcg gggacccca tcagccagc 3300
ccccgttccc ctttccacgt tgccatcagc atcctcggc ttggcaggg accagccgtc 3360
ttccactgcc ttcacccctc tcatatcaac ccgagtgtct cttcgaaaaa cccgccagc 3420
30 tccagagcgg gccagcggcg ccatcaccaa gggcgtggc ttggacagca ccgaggcgct 3480
gtgcctcgcc atctctggga actccgagca gatggccagc cacagcgagc tgctggaggc 3540
cggcaaaaac ctctacacgt tctgctgag ctatgtgat tccatccagc aaatgaggaa 3600
caagtttgcc ttccgagagg ccatcaaca actggagaat aatctccgg agcttcagat 3660
ctgcccggcg tcagcaggca gtggctccgc ggccactcag gacttcagca agctcctcag 3720
35 ttcgggtgaag gaaatcagtg acatagtga gaggtagcag cagtcagggg tcaggtgtca 3780
ggcccgtcgg agctgcctgc agcacatgcg ggctcgcca taccatgac agtggtgag 3840

<210> 4

<211> 1130

<212> PRT

<213> Homo sapiens

5

<220>

<223> Description of Sequence: N/A

<400> 4

10 Met Leu Glu Ile Cys Leu Lys Leu Val Gly Cys Lys Ser Lys Lys Gly
1 5 10 15
Leu Ser Ser Ser Ser Ser Cys Tyr Leu Glu Glu Ala Leu Gln Arg Pro
20 25 30
Val Ala Ser Asp Phe Glu Pro Gln Gly Leu Ser Glu Ala Ala Arg Trp
15 35 40 45
Asn Ser Lys Glu Asn Leu Leu Ala Gly Pro Ser Glu Asn Asp Pro Asn
50 55 60
Leu Phe Val Ala Leu Tyr Asp Phe Val Ala Ser Gly Asp Asn Thr Leu
65 70 75 80
20 Ser Ile Thr Lys Gly Glu Lys Leu Arg Val Leu Gly Tyr Asn His Asn
85 90 95
Gly Glu Trp Cys Glu Ala Gln Thr Lys Asn Gly Gln Gly Trp Val Pro
100 105 110
Ser Asn Tyr Ile Thr Pro Val Asn Ser Leu Glu Lys His Ser Trp Tyr
25 115 120 125
His Gly Pro Val Ser Arg Asn Ala Ala Glu Tyr Pro Leu Ser Ser Gly
130 135 140
Ile Asn Gly Ser Phe Leu Val Arg Glu Ser Glu Ser Ser Pro Ser Gln
145 150 155 160
30 Arg Ser Ile Ser Leu Arg Tyr Glu Gly Arg Val Tyr His Tyr Arg Ile
165 170 175
Asn Thr Ala Ser Asp Gly Lys Leu Tyr Val Ser Ser Glu Ser Arg Phe
180 185 190
Asn Thr Leu Ala Glu Leu Val His His His Ser Thr Val Ala Asp Gly
35 195 200 205
Leu Ile Thr Thr Leu His Tyr Pro Ala Pro Lys Arg Asn Lys Pro Thr
210 215 220
Val Tyr Gly Val Ser Pro Asn Tyr Asp Lys Trp Glu Met Glu Arg Thr

225 230 235 240
 Asp Ile Thr Met Lys His Lys Leu Gly Gly Gly Gln Tyr Gly Glu Val
 245 250 255
 Tyr Glu Gly Val Trp Lys Lys Tyr Ser Leu Thr Val Ala Val Lys Thr
 5 260 265 270
 Leu Lys Glu Asp Thr Met Glu Val Glu Glu Phe Leu Lys Glu Ala Ala
 275 280 285
 Val Met Lys Glu Ile Lys His Pro Asn Leu Val Gln Leu Leu Gly Val
 290 295 300
 10 Cys Thr Arg Glu Pro Pro Phe Tyr Ile Ile Thr Glu Phe Met Thr Tyr
 305 310 315 320
 Gly Asn Leu Leu Asp Tyr Leu Arg Glu Cys Asn Arg Gln Glu Val Asn
 325 330 335
 Ala Val Val Leu Leu Tyr Met Ala Thr Gln Ile Ser Ser Ala Met Glu
 15 340 345 350
 Tyr Leu Glu Lys Lys Asn Phe Ile His Arg Asp Leu Ala Ala Arg Asn
 355 360 365
 Cys Leu Val Gly Glu Asn His Leu Val Lys Val Ala Asp Phe Gly Leu
 370 375 380
 20 Ser Arg Leu Met Thr Gly Asp Thr Tyr Thr Ala His Ala Gly Ala Lys
 385 390 395 400
 Phe Pro Ile Lys Trp Thr Ala Pro Glu Ser Leu Ala Tyr Asn Lys Phe
 405 410 415
 Ser Ile Lys Ser Asp Val Trp Ala Phe Gly Val Leu Leu Trp Glu Ile
 25 420 425 430
 Ala Thr Tyr Gly Met Ser Pro Tyr Pro Gly Ile Asp Arg Ser Gln Val
 435 440 445
 Tyr Glu Leu Leu Glu Lys Asp Tyr Arg Met Lys Arg Pro Glu Gly Cys
 450 455 460
 30 Pro Glu Lys Val Tyr Glu Leu Met Arg Ala Cys Trp Gln Trp Asn Pro
 465 470 475 480
 Ser Asp Arg Pro Ser Phe Ala Glu Ile His Gln Ala Phe Glu Thr Met
 485 490 495
 Phe Gln Glu Ser Ser Ile Ser Asp Glu Val Glu Lys Glu Leu Gly Lys
 35 500 505 510
 Gln Gly Val Arg Gly Ala Val Thr Thr Leu Leu Gln Ala Pro Glu Leu
 515 520 525
 Pro Thr Lys Thr Arg Thr Ser Arg Arg Ala Ala Glu His Arg Asp Thr

	530		535		540	
	Thr Asp Val Pro Glu Met	Pro His Ser Lys Gly Gln Gly Glu Ser Asp				
	545	550	555	560		
	Pro Leu Asp His Glu Pro Ala Val Ser Pro Leu Leu Pro Arg Lys Glu					
5	565	570	575			
	Arg Gly Pro Pro Glu Gly Gly Leu Asn Glu Asp Glu Arg Leu Leu Pro					
	580	585	590			
	Lys Asp Lys Lys Thr Asn Leu Phe Ser Ala Leu Ile Lys Lys Lys Lys					
	595	600	605			
10	Lys Thr Ala Pro Thr Pro Pro Lys Arg Ser Ser Ser Phe Arg Glu Met					
	610	615	620			
	Asp Gly Gln Pro Glu Arg Arg Gly Ala Gly Glu Glu Glu Gly Arg Asp					
	625	630	635	640		
	Ile Ser Asn Gly Ala Leu Ala Phe Thr Pro Leu Asp Thr Ala Asp Pro					
15	645	650	655			
	Ala Lys Ser Pro Lys Pro Ser Asn Gly Ala Gly Val Pro Asn Gly Ala					
	660	665	670			
	Leu Arg Glu Ser Gly Gly Ser Gly Phe Arg Ser Pro His Leu Trp Lys					
	675	680	685			
20	Lys Ser Ser Thr Leu Thr Ser Ser Arg Leu Ala Thr Gly Glu Glu Glu					
	690	695	700			
	Gly Gly Gly Ser Ser Ser Lys Arg Phe Leu Arg Ser Cys Ser Val Ser					
	705	710	715	720		
	Cys Val Pro His Gly Ala Lys Asp Thr Glu Trp Arg Ser Val Thr Leu					
25	725	730	735			
	Pro Arg Asp Leu Gln Ser Thr Gly Arg Gln Phe Asp Ser Ser Thr Phe					
	740	745	750			
	Gly Gly His Lys Ser Glu Lys Pro Ala Leu Pro Arg Lys Arg Ala Gly					
	755	760	765			
30	Glu Asn Arg Ser Asp Gln Val Thr Arg Gly Thr Val Thr Pro Pro Pro					
	770	775	780			
	Arg Leu Val Lys Lys Asn Glu Glu Ala Ala Asp Glu Val Phe Lys Asp					
	785	790	795	800		
	Ile Met Glu Ser Ser Pro Gly Ser Ser Pro Pro Asn Leu Thr Pro Lys					
35	805	810	815			
	Pro Leu Arg Arg Gln Val Thr Val Ala Pro Ala Ser Gly Leu Pro His					
	820	825	830			
	Lys Glu Glu Ala Trp Lys Gly Ser Ala Leu Gly Thr Pro Ala Ala Ala					

	835		840		845
	Glu Pro Val Thr Pro Thr Ser Lys Ala Gly Ser Gly Ala Pro Arg Gly				
	850		855		860
	Thr Ser Lys Gly Pro Ala Glu Glu Ser Arg Val Arg Arg His Lys His				
5	865		870		875
	Ser Ser Glu Ser Pro Gly Arg Asp Lys Gly Lys Leu Ser Lys Leu Lys				
		885		890	895
	Pro Ala Pro Pro Pro Pro Pro Ala Ala Ser Ala Gly Lys Ala Gly Gly				
		900		905	910
10	Lys Pro Ser Gln Arg Pro Gly Gln Glu Ala Ala Gly Glu Ala Val Leu				
		915		920	925
	Gly Ala Lys Thr Lys Ala Thr Ser Leu Val Asp Ala Val Asn Ser Asp				
		930		935	940
	Ala Ala Lys Pro Ser Gln Pro Ala Glu Gly Leu Lys Lys Pro Val Leu				
15	945		950		955
	Pro Ala Thr Pro Lys Pro His Pro Ala Lys Pro Ser Gly Thr Pro Ile				
		965		970	975
	Ser Pro Ala Pro Val Pro Leu Ser Thr Leu Pro Ser Ala Ser Ser Ala				
		980		985	990
20	Leu Ala Gly Asp Gln Pro Ser Ser Thr Ala Phe Ile Pro Leu Ile Ser				
		995		1000	1005
	Thr Arg Val Ser Leu Arg Lys Thr Arg Gln Pro Pro Glu Arg Ala Ser				
		1010		1015	1020
	Gly Ala Ile Thr Lys Gly Val Val Leu Asp Ser Thr Glu Ala Leu Cys				
25	1025		1030		1035
	Leu Ala Ile Ser Gly Asn Ser Glu Gln Met Ala Ser His Ser Ala Val				
		1045		1050	1055
	Leu Glu Ala Gly Lys Asn Leu Tyr Thr Phe Cys Val Ser Tyr Val Asp				
		1060		1065	1070
30	Ser Ile Gln Gln Met Arg Asn Lys Phe Ala Phe Arg Glu Ala Ile Asn				
		1075		1080	1085
	Lys Leu Glu Asn Asn Leu Arg Glu Leu Gln Ile Cys Pro Ala Ser Ala				
		1090		1095	1100
	Gly Ser Gly Pro Ala Ala Thr Gln Asp Phe Ser Lys Leu Leu Ser Ser				
35	1105		1110		1115
	Val Lys Glu Ile Ser Asp Ile Val Gln Arg				
		1125		1130	

<210> 5
<211> 1461
5 <212> DNA
<213> Homo sapiens

<220> MKK7
<223> Description of Sequence: N/A
10

<400> 5
aggcgggtgtt tgtctgccgg actgacgggc ggccggggcgg tgcgcggcgg cggtagggcgg 60
ggggaaaaatg gcggcgctcct ccctggaaca gaagctgtcc cgcttgaag caaagctgaa 120
gcaggagaac cgggaggccc ggcgaggat cgacctcaac ctggatatca gccccagcg 180
15 gcccaggccc accctgcagc tcccgctggc caacgatggg ggcagccgct cgccatcctc 240
agagagctcc ccgcagcacc ccacgcccc cgcccgccc cgccacatgc tggggctccc 300
gtcaaccctg ttcacacccc gcagcatgga gagcattgag attgaccaga agctgcagga 360
gatcatgaag cagacgggct acctgacct cggggggccag cgctaccagg cagaaatcaa 420
cgacctggag aacttgggcy agatgggcag cggcacctgc ggccagggtgt ggaagatgcy 480
20 cttccggaag accggccacg tcattgccgt taagcaaatg cggcgctccg ggaacaagga 540
ggagaacaag cgcacacctca tggacctgga tgtggtgctg aagagccacg actgccccta 600
catcgtgcag tgctttggga cgttcacac caacacggac gtcttcatcg ccatggagct 660
catgggcacc tgcgctgaga agctcaagaa gcggatgcag ggcccatcc ccgagcgcat 720
tctgggcaag atgacagtgg cgattgtgaa ggcgctgtac tacctgaagg agaagcacgg 780
25 tgtcatccac cgcgacgtca agccctcaa catcctgctg gacgagcggg gccagatcaa 840
gttctgcgac ttcggcatca gcggccgcct ggtggactcc aaagccaaga cgcggagcgc 900
cggctgtgcc gcctacatgg caccgagcg cattgacccc ccagaccca ccaagccgga 960
ctatgacatc cgggcccagc tatggagcct gggcatctcg ctggtggagc tggcaacagg 1020
acagtttccc tacaagaact gcaagacgga ctttgaggtc ctcaccaaag tcctacagga 1080
30 agagccccc cttctgcccg gacacatggg cttctcgggg gacttccagt ctttcgtcaa 1140
agactgcctt actaaagatc acaggaagag accaaagtat aataagctac ttgaacacag 1200
cttcatcaag cgctacgaga cgctggaggt ggacgtggcg tcctggttca aggatgtcat 1260
ggcgaagact gagtcaccgc ggactagcgg cgtcctgagc cagccccacc tgcccttctt 1320
caggtagctg cttggcgggc gccagcccca cagggggcca ggggcatggc cacaggcccc 1380
35 cctccccact tggccacca gctgcctgcc aggggagacc tgggacctgg acggccacct 1440
aggactgagg acagagagt g 1461

<210> 6

<211> 419

<212> PRT

<213> Homo sapiens

5

<220>

<223> Description of Sequence: N/A

<400> 6

10 Met Ala Ala Ser Ser Leu Glu Gln Lys Leu Ser Arg Leu Glu Ala Lys
 1 5 10 15
 Leu Lys Gln Glu Asn Arg Glu Ala Arg Arg Arg Ile Asp Leu Asn Leu
 20 25 30
 Asp Ile Ser Pro Gln Arg Pro Arg Pro Thr Leu Gln Leu Pro Leu Ala
 15 35 40 45
 Asn Asp Gly Gly Ser Arg Ser Pro Ser Ser Glu Ser Ser Pro Gln His
 50 55 60
 Pro Thr Pro Pro Ala Arg Pro Arg His Met Leu Gly Leu Pro Ser Thr
 65 70 75 80
 20 Leu Phe Thr Pro Arg Ser Met Glu Ser Ile Glu Ile Asp Gln Lys Leu
 85 90 95
 Gln Glu Ile Met Lys Gln Thr Gly Tyr Leu Thr Ile Gly Gly Gln Arg
 100 105 110
 Tyr Gln Ala Glu Ile Asn Asp Leu Glu Asn Leu Gly Glu Met Gly Ser
 25 115 120 125
 Gly Thr Cys Gly Gln Val Trp Lys Met Arg Phe Arg Lys Thr Gly His
 130 135 140
 Val Ile Ala Val Lys Gln Met Arg Arg Ser Gly Asn Lys Glu Glu Asn
 145 150 155 160
 30 Lys Arg Ile Leu Met Asp Leu Asp Val Val Leu Lys Ser His Asp Cys
 165 170 175
 Pro Tyr Ile Val Gln Cys Phe Gly Thr Phe Ile Thr Asn Thr Asp Val
 180 185 190
 Phe Ile Ala Met Glu Leu Met Gly Thr Cys Ala Glu Lys Leu Lys Lys
 35 195 200 205
 Arg Met Gln Gly Pro Ile Pro Glu Arg Ile Leu Gly Lys Met Thr Val
 210 215 220
 Ala Ile Val Lys Ala Leu Tyr Tyr Leu Lys Glu Lys His Gly Val Ile

225 230 235 240
 His Arg Asp Val Lys Pro Ser Asn Ile L u Leu Asp Glu Arg Gly Gln
 245 250 255
 Ile Lys Phe Cys Asp Phe Gly Ile Ser Gly Arg Leu Val Asp Ser Lys
 5 260 265 270
 Ala Lys Thr Arg Ser Ala Gly Cys Ala Ala Tyr Met Ala Pro Glu Arg
 275 280 285
 Ile Asp Pro Pro Asp Pro Thr Lys Pro Asp Tyr Asp Ile Arg Ala Asp
 290 295 300
 10 Val Trp Ser Leu Gly Ile Ser Leu Val Glu Leu Ala Thr Gly Gln Phe
 305 310 315 320
 Pro Tyr Lys Asn Cys Lys Thr Asp Phe Glu Val Leu Thr Lys Val Leu
 325 330 335
 Gln Glu Glu Pro Pro Leu Leu Pro Gly His Met Gly Phe Ser Gly Asp
 15 340 345 350
 Phe Gln Ser Phe Val Lys Asp Cys Leu Thr Lys Asp His Arg Lys Arg
 355 360 365
 Pro Lys Tyr Asn Lys Leu Leu Glu His Ser Phe Ile Lys Arg Tyr Glu
 370 375 380
 20 Thr Leu Glu Val Asp Val Ala Ser Trp Phe Lys Asp Val Met Ala Lys
 385 390 395 400
 Thr Glu Ser Pro Arg Thr Ser Gly Val Leu Ser Gln Pro His Leu Pro
 405 410 415
 Phe Phe Arg

25

<210> 7

<211> 1050

30

<212> DNA

<213> Homo sapiens

<220> CDC2

<223> Description of Sequence: N/A

35

<400> 7

gggggggggg ggcacttggc ttcaaagctg gctcttggaa attgagcggg gacgagcggc 60
 ttgtttagc tgccgtgcgg ccgcccggga ataataagcc gggatctacc ataccattga 120

```

ctaactatgg aagattatac caaaatagag aaaattggag aagggtaccta tggagttgtg 180
tataagggta gacacaaaac tacagggtcaa gtggttagcca tgaaaaaat cagactagaa 240
agtgaagagg aaggggttcc tagtactgca attcgggaaa tttctctatt aaaggaactt 300
cgtcatccaa atatagtcag tcttcaggat gtgcttatgc aggattccag gttatatctc 360
5 atctttgagt ttctttccat ggatctgaag aaatacttgg attctatccc tcctgggtcag 420
tacatggatt cttcacttgt taagagttat ttataccaaa tcctacaggg gattgtgttt 480
tgtcactcta gaagagttct tcacagagac ttaaaacctc aaaatctctt gattgatgac 540
aaaggaacaa ttaaaactggc tgattttggc cttgccagag cttttggaat acctatcaga 600
gtatatacac atgaggtagt aacactctgg tacagatctc cagaagtatt gctgggggtca 660
10 gctcgttact caactccagt tgacatttgg agtataggca ccatatttgc tgaactagca 720
actaagaaac cacttttcca tggggattca gaaattgatc aactcttcag gattttcaga 780
gctttgggca ctcccaataa tgaagtgtgg ccagaagtgg aatctttaca ggactataag 840
aatacatttc ccaaattgaa accaggaagc ctagcatccc atgtcaaaaa cttggatgaa 900
aatggcttgg atttgctctc gaaaatgtta atctatgatc cagccaaacg aatttctggc 960
15 aaaatggcac tgaatcatcc atattttaat gatttggaca atcagattaa gaagatgtag 1020
ctttctgaca aaaagtttcc atatgttatg 1050

```

<210> 8

20 <211> 297

<212> PRT

<213> Homo sapiens

<220>

25 <223> Description of Sequence: N/A

<400> 8

```

Met Glu Asp Tyr Thr Lys Ile Glu Lys Ile Gly Glu Gly Thr Tyr Gly
  1             5             10             15
30 Val Val Tyr Lys Gly Arg His Lys Thr Thr Gly Gln Val Val Ala Met
      20             25             30
Lys Lys Ile Arg Leu Glu Ser Glu Glu Glu Gly Val Pro Ser Thr Ala
      35             40             45
Ile Arg Glu Ile Ser Leu Leu Lys Glu Leu Arg His Pro Asn Ile Val
35      50             55             60
Ser Leu Gln Asp Val Leu Met Gln Asp Ser Arg Leu Tyr Leu Ile Phe
      65             70             75             80
Glu Phe Leu Ser Met Asp Leu Lys Lys Tyr Leu Asp Ser Ile Pro Pro

```

85 90 95
 Gly Gln Tyr Met Asp Ser Ser Leu Val Lys Ser Tyr Leu Tyr Gln Ile
 100 105 110
 Leu Gln Gly Ile Val Phe Cys His Ser Arg Arg Val Leu His Arg Asp
 5 115 120 125
 Leu Lys Pro Gln Asn Leu Leu Ile Asp Asp Lys Gly Thr Ile Lys Leu
 130 135 140
 Ala Asp Phe Gly Leu Ala Arg Ala Phe Gly Ile Pro Ile Arg Val Tyr
 145 150 155 160
 10 Thr His Glu Val Val Thr Leu Trp Tyr Arg Ser Pro Glu Val Leu Leu
 165 170 175
 Gly Ser Ala Arg Tyr Ser Thr Pro Val Asp Ile Trp Ser Ile Gly Thr
 180 185 190
 Ile Phe Ala Glu Leu Ala Thr Lys Lys Pro Leu Phe His Gly Asp Ser
 15 195 200 205
 Glu Ile Asp Gln Leu Phe Arg Ile Phe Arg Ala Leu Gly Thr Pro Asn
 210 215 220
 Asn Glu Val Trp Pro Glu Val Glu Ser Leu Gln Asp Tyr Lys Asn Thr
 225 230 235 240
 20 Phe Pro Lys Trp Lys Pro Gly Ser Leu Ala Ser His Val Lys Asn Leu
 245 250 255
 Asp Glu Asn Gly Leu Asp Leu Leu Ser Lys Met Leu Ile Tyr Asp Pro
 260 265 270
 Ala Lys Arg Ile Ser Gly Lys Met Ala Leu Asn His Pro Tyr Phe Asn
 25 275 280 285
 Asp Leu Asp Asn Gln Ile Lys Lys Met
 290 295

30

<210> 9

<211> 1480

<212> DNA

<213> Homo sapiens

35

<220> CamKI

<223> Description of Sequence: N/A

<400> 9

gaattccgag caagagcgcg ggccgggtggc ccaggcacgc agcgggtgag gaccgcgccc 60
acagctcggc gcccaaccacc gcgggcctcc cagccagccc cgcnnngagc cgcaggancc 120
ctggctgtgg tcggggggca gtgggccatg ctgggggcag tggaaggccc cagggtggaag 180
5 caggcggagg acattagaga catctacgac ttccgagatg ttctgggcac gggggccttc 240
tcggaggtga tcctggcaga agataagagg acgcagaagc tgggtggccat caaatgcatt 300
gcccaaggagg ccctggaggg caaggaaggc agcatggaga atgagattgc tgtcctgcac 360
aagatcaagc accccaacat tgtagccctg gatgacatct atgagagtgg gggccacctc 420
tacctcatca tgcagctggt gtccgggtggg gagctctttg accgtattgt ggaaaaaggc 480
10 ttctacacgg agcgggacgc cagccgcctc atcttcagg tgctggatgc tgtgaaatac 540
ctgcatgacc tgggcattgt acaccgggat ctcaagccag agaatctgct gtactacagc 600
ctggatgaag actccaaaat catgatctcc gactttggcc tctccaagat ggaggaccgc 660
ggcagtgtgc tctccaccgc ctgtggaact ccgggatacg tggcccctga agtcctggcc 720
cagaagccct acagcaaggc tgtggattgc tgggtccatag gtgtcatcgc ctacatcttg 780
15 ctctgcgggt accctccctt ctatgacgag aatgatgcca aactctttga acagatcttg 840
aaggccgagt acgagtttga ctctccttac tgggacgaca tctctgactc tgccaaagat 900
ttcatccggc acttgatgga gaaggacca gagaaaagat tcacctgtga gcaggccttg 960
cagcaccat ggattgcagg agatacagct ctagataaga atatccacca gtcggtgagt 1020
gagcagatca agaagaactt tgccaagagc aagtggaagc aagccttcaa tgccacggct 1080
20 gtggtgctgc acatgaggaa actgcagctg ggcaccagcc aggaggggca ggggcagacg 1140
gcgagccatg gggagctgct gacaccagtg gctggggggc cggcagctgg ctgttgctgt 1200
cgagactgct gcgtggagcc gggcacagaa ctgtccccc cactgcccc ccagctctag 1260
ggccctggac ctccgggtcat gatcctctgc gtgggagggc ttgggggcca gcctgctccc 1320
cttccctccc tgaaccggga gtttctctgc cctgtccct cctcacctgc ttccctacca 1380
25 ctctcactg cattttccat acaaagtgtt ctattttatt gttccttctt gtaataaagg 1440
gaagataaaa ccaaaaaaaaa aaaaaaaaaa acggaattcc 1480

<210> 10

30 <211> 370

<212> PRT

<213> Homo sapiens

<220>

35 <223> Description of Sequence: N/A

<400> 10

Met Leu Gly Ala Val Glu Gly Pro Arg Trp Lys Gln Ala Glu Asp Ile

	1	5	10	15
	Arg	Asp	Ile	Tyr
	Asp	Phe	Arg	Asp
	Val	Leu	Gly	Thr
	Gly	Ala	Phe	Ser
	20	25	30	
	Glu	Val	Ile	Leu
	Ala	Glu	Asp	Lys
	Arg	Thr	Gln	Lys
	Leu	Val	Ala	Ile
5	35	40	45	
	Lys	Cys	Ile	Ala
	Lys	Glu	Ala	Leu
	Glu	Gly	Lys	Glu
	Gly	Ser	Met	Glu
	50	55	60	
	Asn	Glu	Ile	Ala
	Val	Leu	His	Lys
	Ile	Lys	His	Pro
	Asn	Ile	Val	Ala
	65	70	75	80
10	Leu	Asp	Asp	Ile
	Tyr	Glu	Ser	Gly
	Gly	Gly	His	Leu
	Tyr	Leu	Ile	Met
	Gln			
	85	90	95	
	Leu	Val	Ser	Gly
	Gly	Glu	Leu	Phe
	Asp	Arg	Ile	Val
	Glu	Lys	Gly	Phe
	100	105	110	
	Tyr	Thr	Glu	Arg
	Asp	Ala	Ser	Arg
	Leu	Ile	Phe	Gln
	Val	Leu	Asp	Ala
15	115	120	125	
	Val	Lys	Tyr	Leu
	His	Asp	Leu	Gly
	Ile	Val	His	Arg
	Asp	Leu	Lys	Pro
	130	135	140	
	Glu	Asn	Leu	Leu
	Tyr	Tyr	Ser	Leu
	Asp	Glu	Asp	Ser
	Lys	Ile	Met	Ile
	145	150	155	160
20	Ser	Asp	Phe	Gly
	Leu	Ser	Lys	Met
	Glu	Asp	Pro	Gly
	Ser	Val	Leu	Ser
	165	170	175	
	Thr	Ala	Cys	Gly
	Thr	Pro	Gly	Tyr
	Val	Ala	Pro	Glu
	Val	Leu	Ala	Gln
	180	185	190	
	Lys	Pro	Tyr	Ser
	Lys	Ala	Val	Asp
	Cys	Trp	Ser	Ile
	Gly	Val	Ile	Ala
25	195	200	205	
	Tyr	Ile	Leu	Leu
	Cys	Gly	Tyr	Pro
	Pro	Phe	Tyr	Asp
	Glu	Asn	Asp	Ala
	210	215	220	
	Lys	Leu	Phe	Glu
	Gln	Ile	Leu	Lys
	Ala	Glu	Tyr	Glu
	Phe	Asp	Ser	Pro
	225	230	235	240
30	Tyr	Trp	Asp	Asp
	Ile	Ser	Asp	Ser
	Ala	Lys	Asp	Phe
	Ile	Arg	His	Leu
	245	250	255	
	Met	Glu	Lys	Asp
	Pro	Glu	Lys	Arg
	Phe	Thr	Cys	Glu
	Gln	Ala	Leu	Gln
	260	265	270	
	His	Pro	Trp	Ile
	Ala	Gly	Asp	Thr
	Ala	Leu	Asp	Lys
	Asn	Ile	His	Gln
35	275	280	285	
	Ser	Val	Ser	Glu
	Gln	Ile	Lys	Lys
	Asn	Phe	Ala	Lys
	Ser	Lys	Trp	Lys
	290	295	300	
	Gln	Ala	Phe	Asn
	Ala	Thr	Ala	Val
	Val	Arg	His	Met
	Arg	Lys	Leu	Gln

10

<210> 11

<211> 1782

15

<212> DNA

<213> Homo sapiens

<220> JNK2

<223> Description of Sequence: N/A

20

<400> 11

gggcggggcga	gggatctgaa	acttgcccac	ccttcgggat	attgcaggac	gctgcatcat	60
gagcgacagt	aaatgtgaca	gtcagtttta	tagtgtgcaa	gtggcagact	caaccttcac	120
tgtcctaata	cgttaccagc	agctgaaacc	aattggctct	ggggcccaag	ggattgtttg	180
tgtctgcat	gatacagttc	ttgggataag	tgttgcaagc	aagaaactaa	gccgtccttt	240
tcagaaccaa	actcatgcaa	agagagctta	tcgtgaactt	gtcctcttaa	aatgtgtcaa	300
tcataaaaa	ataattagtt	tgttaaatgt	gtttacacca	caaaaaactc	tagaagaatt	360
tcaagatgtg	tatttggtta	tggaaatta	ggatgctaac	ttatgtcagg	ttattcacat	420
ggagctggat	catgaaagaa	tgtcctacct	tctttaccag	atgctttgtg	gtattaaaca	480
tctgcattca	gctgggtata	ttcatagaga	tttgaagcct	agcaacattg	ttgtgaaatc	540
agactgcacc	ctgaagatcc	ttgactttgg	cctggcccg	acagcgtgca	ctaacttcat	600
gatgaccct	tacgtggtga	cacgggtacta	ccgggcgccc	gaagtcatcc	tgggtatggg	660
ctacaaagag	aacgttgata	tctgggtcagt	gggttgcatc	atgggagagc	tgggtgaaagg	720
ttgtgtgata	ttccaaggca	ctgaccatat	tgatcagtgg	aataaagtta	ttgagcagct	780
gggaacacca	tcagcagagt	tcatgaagaa	acttcagcca	actgtgagga	attatgtcga	840
aaacagacca	aagtatcctg	gaatcaaatt	tgaagaactc	tttccagatt	ggatattccc	900
atcagaatct	gagcgagaca	aaataaaaac	aagtcaagcc	agagatctgt	tatcaaaaat	960
gttagtgatt	gatacctgaca	agcggatctc	tgtagacgaa	gctctgcgtc	accatacat	1020

cactgtttgg tatgaccccg ccgaagcaga agccccacca cctcaaattt atgatgccca 1080
 gttggaagaa agagaacatg caattgaaga atggaaagag ctaatttaca aagaagtcac 1140
 ggattgggaa gaaagaagca agaattggtg tgtaaaagat cagccttcag atgcagcagt 1200
 aagtagcaac gccactcctt ctgagtcctt atcgatcaat gacatttcat ccatgtccac 1260
 5 tgagcagacg ctggcctcag acacagacag cagtcttgat gcctcgacgg gacccttga 1320
 aggctgtcga tgataggta gaaatagcaa acctgtcagc attgaaggaa ctctcacctc 1380
 cgtgggcctg aaatgcttgg gagttgatgg aaccaaatag aaaaactcca tgttctgcat 1440
 gtaagaaaca caatgccttg ccctattcag acctgatagg attgcctgct tagatgataa 1500
 aatgaggcag aatatgtctg aagaaaaaaa ttgcaagcca cacttctaga gattttgttc 1560
 10 aagatcattt cagggtgagca gttagagtag gtgaatttgt ttcaaattgt actagtgaca 1620
 gtttctcatc atctgtaact gttgagatgt atgtgcatgt gaccacaaat gcttgcttgg 1680
 acttgcccat ctagcacttt ggaaatcagt atttaaagtc caaataatct tccaggtagt 1740
 gctgcttctg aagttatctc ttaatcctct taagtaattt gg 1782

15

<210> 12

<211> 424

<212> PRT

<213> Homo sapiens

20

<220>

<223> Description of Sequence: N/A

<400> 12

25

Met Ser Asp Ser Lys Cys Asp Ser Gln Phe Tyr Ser Val Gln Val Ala

1

5

10

15

Asp Ser Thr Phe Thr Val Leu Lys Arg Tyr Gln Gln Leu Lys Pro Ile

20

25

30

Gly Ser Gly Ala Gln Gly Ile Val Cys Ala Ala Phe Asp Thr Val Leu

30

35

40

45

Gly Ile Ser Val Ala Val Lys Lys Leu Ser Arg Pro Phe Gln Asn Gln

50

55

60

Thr His Ala Lys Arg Ala Tyr Arg Glu Leu Val Leu Leu Lys Cys Val

65

70

75

80

35

Asn His Lys Asn Ile Ile Ser Leu Leu Asn Val Phe Thr Pro Gln Lys

85

90

95

Thr Leu Glu Glu Phe Gln Asp Val Tyr Leu Val Met Glu Leu Met Asp

100

105

110

Ala Asn Leu Cys Gln Val Ile His Met Glu Leu Asp His Glu Arg Met
 115 120 125
 Ser Tyr Leu Leu Tyr Gln Met Leu Cys Gly Ile Lys His Leu His Ser
 130 135 140
 5 Ala Gly Ile Ile His Arg Asp Leu Lys Pro Ser Asn Ile Val Val Lys
 145 150 155 160
 Ser Asp Cys Thr Leu Lys Ile Leu Asp Phe Gly Leu Ala Arg Thr Ala
 165 170 175
 Cys Thr Asn Phe Met Met Thr Pro Tyr Val Val Thr Arg Tyr Tyr Arg
 10 180 185 190
 Ala Pro Glu Val Ile Leu Gly Met Gly Tyr Lys Glu Asn Val Asp Ile
 195 200 205
 Trp Ser Val Gly Cys Ile Met Gly Glu Leu Val Lys Gly Cys Val Ile
 210 215 220
 15 Phe Gln Gly Thr Asp His Ile Asp Gln Trp Asn Lys Val Ile Glu Gln
 225 230 235 240
 Leu Gly Thr Pro Ser Ala Glu Phe Met Lys Lys Leu Gln Pro Thr Val
 245 250 255
 Arg Asn Tyr Val Glu Asn Arg Pro Lys Tyr Pro Gly Ile Lys Phe Glu
 20 260 265 270
 Glu Leu Phe Pro Asp Trp Ile Phe Pro Ser Glu Ser Glu Arg Asp Lys
 275 280 285
 Ile Lys Thr Ser Gln Ala Arg Asp Leu Leu Ser Lys Met Leu Val Ile
 290 295 300
 25 Asp Pro Asp Lys Arg Ile Ser Val Asp Glu Ala Leu Arg His Pro Tyr
 305 310 315 320
 Ile Thr Val Trp Tyr Asp Pro Ala Glu Ala Glu Ala Pro Pro Pro Gln
 325 330 335
 Ile Tyr Asp Ala Gln Leu Glu Glu Arg Glu His Ala Ile Glu Glu Trp
 30 340 345 350
 Lys Glu Leu Ile Tyr Lys Glu Val Met Asp Trp Glu Glu Arg Ser Lys
 355 360 365
 Asn Gly Val Val Lys Asp Gln Pro Ser Asp Ala Ala Val Ser Ser Asn
 370 375 380
 35 Ala Thr Pro Ser Gln Ser Ser Ser Ile Asn Asp Ile Ser Ser Met Ser
 385 390 395 400
 Thr Glu Gln Thr Leu Ala Ser Asp Thr Asp Ser Ser Leu Asp Ala Ser
 405 410 415

Thr Gly Pro Leu Glu Gly Cys Arg

420

5

<210> 13

<211> 3668

<212> DNA

<213> Homo sapiens

10

<220> LIMK-2

<223> Description of Sequence: N/A

<400> 13

15 gtggtcttcc cgcgcctgag gcggcggcgg caggagctga ggggagttgt agggaaactga 60
ggggagctgc tgtgtccccc gcctcctcct cccatttcc gggctcccgg gaccatgtcc 120
gcgctggcgg gtgaagatgt ctggaggtgt ccaggctgtg gggaccacat tgctccaagc 180
cagatatggt acaggactgt caacgaaacc tggcacggct cttgcttccg gtgttcagaa 240
tgccaggatt ccctcaccaa ctggtactat gagaaggatg ggaagctcta ctgccccaa 300
20 gactactggg ggaagtttgg ggagttctgt catgggtgct ccctgctgat gacagggcct 360
tttatggtgg ctggggagtt caagtaccac ccagagtgt ttgctgtat gagctgcaag 420
gtgatcattg aggatgggga tgcataatgca ctggtgcagc atgccaccct ctactgtggg 480
aagtgccaca atgagtggtg gctggcacc atgtttgaga gactctccac agagtctgtt 540
caggagcagc tgccctactc tgtcacgtc atctccatgc cggccaccac tgaaggcagg 600
25 cggggcttct ccgtgtccgt ggagagtgcc tgcctcaact acgccaccac tgtgcaagt 660
aaagaggtca accggtatgca catcagtccc aacaatcgaa acgccatcca ccctggggac 720
cgcacctctg agatcaatgg gacccccgtc cgcacacttc gaggggagga ggtggaggat 780
gcaattagcc agacgagcca gacacttcag ctgttgattg aacatgacc cgtctcccaa 840
cgcttgacc agctgcggct ggaggcccg ctcgtctctc acatgcagaa tgccggacac 900
30 cccacgccc tcagcacctt ggacaccaag gagaatctgg aggggacact gaggagacgt 960
tccctaagge gcagtaacag tatctccaag tccctggcc ccagctccc aaaggagccc 1020
ctgctgttca gccgtgacat cagccgtcga gaatcccttc gttgttccag cagctattca 1080
cagcagatct tccggccctg tgacctaatc catggggagg tcctggggaa gggcttcttt 1140
gggcaggcta tcaaggtgac acacaaagcc acgggcaaag tgatggtcat gaaagagtta 1200
35 attcgatgtg atgaggagac ccagaaaact tttctgactg aggtgaaagt gatgcgcagc 1260
ctggaccacc ccaatgtgct caagttcatt ggtgtgctgt acaaggataa gaagctgaac 1320
ctgctgacag agtacattga ggggggcaca ctgaaggact ttctgcgag tatggatccg 1380
ttcccctggc agcagaaggt caggtttgcc aaaggaatcg cctccggaat ggcctatttg 1440

cactctatgt gcatcatcca ccgggatctg aactcgacaca actgcctcat caagttggac 1500
aagactgtgg tggtaggcaga ctttgggctg tcacggctca tagtggaaga gaggaaaagg 1560
gcccccatgg agaaggccac caccaagaaa cgcaccttgc gcaagaacga ccgcaagaag 1620
cgctacacgg tggtagggaaa ccctactggt atggccctg agatgctgaa cggaaagagc 1680
5 tatgatgaga cggtaggat cttctccttt gggatcggtt tctgtgagat cattgggcag 1740
gtgtatgcag atcctgactg ccttccccga acactggact ttggcctcaa cgtgaagctt 1800
ttctgggaga agtttgttcc cacagattgt ccccgccct tcttcccgct ggccgccatc 1860
tgctgcagac tggagcctga gagcagacca gcattctcga aattggagga ctcttttgag 1920
gccctctccc tgtacctggg ggagctgggc atcccgtgc ctgcagagct ggaggagttg 1980
10 gaccacactg tgagcatgca gtacggcctg acccgggact cacctcccta gccctggccc 2040
agccccctgc aggggggtgt tctacagcca gcattgcccc tctgtgcccc attcctgctg 2100
tgagcagggc cgtccgggct tctgtggat tggcggaatg tttagaagca gaacaagcca 2160
ttcctattac ctccccagga ggcaagtggg cgcagcacca gggaaatgta tctccacagg 2220
ttctggggcc tagttactgt ctgtaaactc aatacttgcc tgaaagctgt gaagaagaaa 2280
15 aaaaccctg gcctttgggc caggaggaat ctgttactg aatccacca ggaactccct 2340
ggcagtggat tgtgggagc tcttgcttac actaatcagc gtgacctgga cctgctgggc 2400
aggatcccag ggtgaacctg cctgtgaact ctgaagtcac tagtccagct ggggtgcagga 2460
ggacttcaag tgtgtggacg aaagaaagac tgatggctca aagggtgtga aaaagtcagt 2520
gatgctcccc ctttctactc cagatcctgt cttcctgga gcaaggttga gggagtaggt 2580
20 tttgaagagt cccttaatat gtggtggaac aggccaggag ttagagaaaag ggctggcttc 2640
tgtttacctg ctactggct ctagccagcc cagggaccac atcaatgtga gaggaagcct 2700
ccacctcatg ttttcaaact taatactgga gactggctga gaacttacgg acaacatcct 2760
ttctgtctga acaaacagt cacaagcaca ggaagaggct gggggactag aaagaggccc 2820
tgccctctag aaagctcaga tcttggttc tgttactcat actcgggtgg gctccttagt 2880
25 cagatgccta aaacattttg cctaaagctc gatgggttct ggaggacagt gtggcttgtc 2940
acaggcctag agtctgaggg aggggagtg gagtctcagc aatctcttgg tcttggttc 3000
atggcaacca ctgctacccc ttcaacatgc ctggtttagg cagcagcttg ggctgggaag 3060
aggtgggtgg agagtctcaa agctgagatg ctgagagaga tagctccctg agctgggcca 3120
tctgacttct acctccatg tttgctctcc caactcatta gctcctgggc agcatcctcc 3180
30 tgagccacat gtgcaggtac tggaaaacct ccatcttggc tcccagagct ctaggaactc 3240
ttcatcaca ctagatttgc ctcttctaag tgtctatgag cttgcacat atttaataaa 3300
ttgggaatgg gtttgggta ttaatgcaat gtgtggtggt tgtattggag cagggggaat 3360
tgataaagga gagtgggtgc tgttaatat atcttatcta ttgggtggt tgtgaaatat 3420
tgtacataga cctgatgagt tgtgggacca gatgtcatct ctggtcagag tttacttgct 3480
35 atatagactg tacttatgtg tgaagtttgc aagcttgctt tagggctgag ccctggactc 3540
ccagcagcag cacagttcag cattgtgtgg ctggttgttt cctggctgtc ccagcaagt 3600
gtaggagtgg tgggcctgaa ctgggccatt gatcagacta aataaattaa gcagttaaca 3660
taactggc 3668

<210> 14
 <211> 638
 5 <212> PRT
 <213> Homo sapiens

 <220>
 <223> Description of Sequence: N/A
 10
 <400> 14
 Met Ser Ala Leu Ala Gly Glu Asp Val Trp Arg Cys Pro Gly Cys Gly
 1 5 10 15
 Asp His Ile Ala Pro Ser Gln Ile Trp Tyr Arg Thr Val Asn Glu Thr
 15 20 25 30
 Trp His Gly Ser Cys Phe Arg Cys Ser Glu Cys Gln Asp Ser Leu Thr
 35 40 45
 Asn Trp Tyr Tyr Glu Lys Asp Gly Lys Leu Tyr Cys Pro Lys Asp Tyr
 50 55 60
 20 Trp Gly Lys Phe Gly Glu Phe Cys His Gly Cys Ser Leu Leu Met Thr
 65 70 75 80
 Gly Pro Phe Met Val Ala Gly Glu Phe Lys Tyr His Pro Glu Cys Phe
 85 90 95
 Ala Cys Met Ser Cys Lys Val Ile Ile Glu Asp Gly Asp Ala Tyr Ala
 25 100 105 110
 Leu Val Gln His Ala Thr Leu Tyr Cys Gly Lys Cys His Asn Glu Val
 115 120 125
 Val Leu Ala Pro Met Phe Glu Arg Leu Ser Thr Glu Ser Val Gln Glu
 130 135 140
 30 Gln Leu Pro Tyr Ser Val Thr Leu Ile Ser Met Pro Ala Thr Thr Glu
 145 150 155 160
 Gly Arg Arg Gly Phe Ser Val Ser Val Glu Ser Ala Cys Ser Asn Tyr
 165 170 175
 Ala Thr Thr Val Gln Val Lys Glu Val Asn Arg Met His Ile Ser Pro
 35 180 185 190
 Asn Asn Arg Asn Ala Ile His Pro Gly Asp Arg Ile Leu Glu Ile Asn
 195 200 205
 Gly Thr Pro Val Arg Thr Leu Arg Val Glu Glu Val Glu Asp Ala Ile

	210		215		220	
	Ser Gln Thr Ser Gln Thr	Leu Gln Leu Leu Ile Glu His Asp Pro Val				
	225	230	235	240		
	Ser Gln Arg Leu Asp Gln Leu Arg Leu Glu Ala Arg Leu Ala Pro His					
5	245	250	255			
	Met Gln Asn Ala Gly His Pro His Ala Leu Ser Thr Leu Asp Thr Lys					
	260	265	270			
	Glu Asn Leu Glu Gly Thr Leu Arg Arg Arg Ser Leu Arg Arg Ser Asn					
	275	280	285			
10	Ser Ile Ser Lys Ser Pro Gly Pro Ser Ser Pro Lys Glu Pro Leu Leu					
	290	295	300			
	Phe Ser Arg Asp Ile Ser Arg Ser Glu Ser Leu Arg Cys Ser Ser Ser					
	305	310	315	320		
	Tyr Ser Gln Gln Ile Phe Arg Pro Cys Asp Leu Ile His Gly Glu Val					
15	325	330	335			
	Leu Gly Lys Gly Phe Phe Gly Gln Ala Ile Lys Val Thr His Lys Ala					
	340	345	350			
	Thr Gly Lys Val Met Val Met Lys Glu Leu Ile Arg Cys Asp Glu Glu					
	355	360	365			
20	Thr Gln Lys Thr Phe Leu Thr Glu Val Lys Val Met Arg Ser Leu Asp					
	370	375	380			
	His Pro Asn Val Leu Lys Phe Ile Gly Val Leu Tyr Lys Asp Lys Lys					
	385	390	395	400		
	Leu Asn Leu Leu Thr Glu Tyr Ile Glu Gly Gly Thr Leu Lys Asp Phe					
25	405	410	415			
	Leu Arg Ser Met Asp Pro Phe Pro Trp Gln Gln Lys Val Arg Phe Ala					
	420	425	430			
	Lys Gly Ile Ala Ser Gly Met Ala Tyr Leu His Ser Met Cys Ile Ile					
	435	440	445			
30	His Arg Asp Leu Asn Ser His Asn Cys Leu Ile Lys Leu Asp Lys Thr					
	450	455	460			
	Val Val Val Ala Asp Phe Gly Leu Ser Arg Leu Ile Val Glu Glu Arg					
	465	470	475	480		
	Lys Arg Ala Pro Met Glu Lys Ala Thr Thr Lys Lys Arg Thr Leu Arg					
35	485	490	495			
	Lys Asn Asp Arg Lys Lys Arg Tyr Thr Val Val Gly Asn Pro Tyr Trp					
	500	505	510			
	Met Ala Pro Glu Met Leu Asn Gly Lys Ser Tyr Asp Glu Thr Val Asp					

```

          515              520              525
Ile Phe Ser Phe Gly Ile Val Leu Cys Glu Ile Ile Gly Gln Val Tyr
          530              535              540
Ala Asp Pro Asp Cys Leu Pro Arg Thr Leu Asp Phe Gly Leu Asn Val
5  545              550              555              560
Lys Leu Phe Trp Glu Lys Phe Val Pro Thr Asp Cys Pro Pro Ala Phe
          565              570              575
Phe Pro Leu Ala Ala Ile Cys Cys Arg Leu Glu Pro Glu Ser Arg Pro
          580              585              590
10 Ala Phe Ser Lys Leu Glu Asp Ser Phe Glu Ala Leu Ser Leu Tyr Leu
          595              600              605
Gly Glu Leu Gly Ile Pro Leu Pro Ala Glu Leu Glu Glu Leu Asp His
          610              615              620
Thr Val Ser Met Gln Tyr Gly Leu Thr Arg Asp Ser Pro Pro
15 625              630              635

```

```

<210> 15
20 <211> 2169
   <212> DNA
   <213> Homo sapiens

```

```

<220> PRK
25 <223> Description of Sequence: N/A

```

```

<400> 15
ccgcctccga gtgccttgcg cggacctgag ctggagatgc tggccgggct accgacgtca 60
gaccccgggc gcctcatcac ggaccgcgc agcggccgca cctacctcaa aggccgcttg 120
30 ttgggcaagg ggggcttcgc ccgctgctac gaggccactg acacagagac tggcagcgcc 180
tacgctgtca aagtcatccc gcagagccgc gtcgccaagc cgcacagcg cgagaagatc 240
ctaaatgaga ttgagctgca ccgagacctg cagcaccgcc acatcgctgcg tttttcgcac 300
cactttgagg acgctgacaa catctacatt ttcttgagc tctgcagccg aaagtccctg 360
gcccacatct ggaaggcccg gcacaccctg ttggagccag aagtgcgcta ctacctgcgg 420
35 cagatccttt ctggcctcaa gtacttgca cagcgcgga tcttgaccg ggacctcaag 480
ttgggaaatt ttttcatcac tgagaacatg gaactgaagg tgggggattt tgggctggca 540
gcccggttgg agcctccgga gcagaggaag aagaccatct gtggcaccac caactatgtg 600
gctccagaag tgctgctgag acagggccac ggcctgaag cggatgtatg gtcactgggc 660

```

```

tgtgtcatgt acacgctgct ctgcgggagc cctccctttg agacggctga cctgaaggag 720
acgtaccgct gcatcaagca gggtcactac acgtgcctg ccagcctctc actgcctgcc 780
cggcagctcc tggccgccat ccttcgggac tcaccccgag accgcccctc tattgaccag 840
atcctgcgcc atgacttctt taccaagggc tacaccccg atcgactccc tatcagcagc 900
5  tgcgtgacag tcccagacct gacaccccc aaccagcta ggagtctgtt tgccaaagtt 960
accaagagcc tctttggcag aaagaagaag agtaagaatc atgcccagga gagggatgag 1020
gtctccgggt tggtgagcgg cctcatgcgc acatccgttg gccatcagga tgccaggcca 1080
gaggetccag cagcttcttg cccagcccct gtcagcctgg tagagacagc acctgaagac 1140
agctcacccc gtgggacact ggcaagcagt ggagatggat ttgaagaagg tctgactgtg 1200
10 gccacagtag tggagtcagc cctttgtgct ctgagaaatt gtatagcttt catgccccca 1260
gcggaacaga acccgcccc cctggcccag ccagagcctc tgggtgtgggt cagcaagtgg 1320
gttgactact ccaataagtt cggctttggg tatcaactgt ccagccgccg tgtggtgtg 1380
ctcttcaacg atggcacaca tatggccctg tcggccaaca gaaagactgt gcactacaat 1440
cccaccagca caaagcactt ctcttctcc gtgggtgctg tgccccgggc cctgcagcct 1500
15 cagctgggta tctgcggta cttgcctcc tacatggagc agcacctcat gaagggtgga 1560
gatctgcca gtgtggaaga ggtagagga cctgctccgc ccttgtgct gcagtgggtc 1620
aagacggatc aggtctctct catgctgttt agtgatggca ctgtccaggt gaacttctac 1680
ggggaccaca ccaagctgat tctcagtggc tgggagcccc tccttgtgac ttttgtggc 1740
cgaaatcgta gtgcttgta ttacctcgt tccaccttc ggcagctggg ctgctctcca 1800
20 gacctgcggc agcgactccg ctatgctctg cgcctgctcc gggaccgcag cccagcttag 1860
gacccaagcc ctgaaggcct gaggcctgtg cctgtcagga tctggccctt gcctttgtgg 1920
ccttccccct tcttttggtg cctcactggg ggctttgggc cgaatcccc agggaatcag 1980
ggaccagctt tactggagt gggtggcgct tgtcttcgct ggctcctacc ccatctccaa 2040
gataagcctg agccttagct cccagctagg gggcgttatt tatggaccac ttttatttat 2100
25 tgtcagacac ttatttattg ,ggatgtgagc cccagggggc ctctcctag gataataaac 2160
aattttgca 2169

```

<210> 16

30 <211> 607

<212> PRT

<213> Homo sapiens

<220>

35 <223> Description of Sequence: N/A

<400> 16

Met Leu Ala Gly Leu Pro Thr Ser Asp Pro Gly Arg Leu Ile Thr Asp

	305		310		315		320									
	Asn	His	Ala	Gln	Glu	Arg	Asp	Glu	Val	Ser	Gly	Leu	Val	Ser	Gly	Leu
				325						330					335	
	Met	Arg	Thr	Ser	Val	Gly	His	Gln	Asp	Ala	Arg	Pro	Glu	Ala	Pro	Ala
5				340						345					350	
	Ala	Ser	Gly	Pro	Ala	Pro	Val	Ser	Leu	Val	Glu	Thr	Ala	Pro	Glu	Asp
				355						360					365	
	Ser	Ser	Pro	Arg	Gly	Thr	Leu	Ala	Ser	Ser	Gly	Asp	Gly	Phe	Glu	Glu
				370						375					380	
10	Gly	Leu	Thr	Val	Ala	Thr	Val	Val	Glu	Ser	Ala	Leu	Cys	Ala	Leu	Arg
				385						390					395	
	Asn	Cys	Ile	Ala	Phe	Met	Pro	Pro	Ala	Glu	Gln	Asn	Pro	Ala	Pro	Leu
				405						410					415	
	Ala	Gln	Pro	Glu	Pro	Leu	Val	Trp	Val	Ser	Lys	Trp	Val	Asp	Tyr	Ser
15				420						425					430	
	Asn	Lys	Phe	Gly	Phe	Gly	Tyr	Gln	Leu	Ser	Ser	Arg	Arg	Val	Ala	Val
				435						440					445	
	Leu	Phe	Asn	Asp	Gly	Thr	His	Met	Ala	Leu	Ser	Ala	Asn	Arg	Lys	Thr
				450						455					460	
20	Val	His	Tyr	Asn	Pro	Thr	Ser	Thr	Lys	His	Phe	Ser	Phe	Ser	Val	Gly
				465						470					475	
	Ala	Val	Pro	Arg	Ala	Leu	Gln	Pro	Gln	Leu	Gly	Ile	Leu	Arg	Tyr	Phe
				485						490					495	
	Ala	Ser	Tyr	Met	Glu	Gln	His	Leu	Met	Lys	Gly	Gly	Asp	Leu	Pro	Ser
25				500						505					510	
	Val	Glu	Glu	Val	Glu	Val	Pro	Ala	Pro	Pro	Leu	Leu	Leu	Gln	Trp	Val
				515						520					525	
	Lys	Thr	Asp	Gln	Ala	Leu	Leu	Met	Leu	Phe	Ser	Asp	Gly	Thr	Val	Gln
				530						535					540	
30	Val	Asn	Phe	Tyr	Gly	Asp	His	Thr	Lys	Leu	Ile	Leu	Ser	Gly	Trp	Glu
				545						550					555	
	Pro	Leu	Leu	Val	Thr	Phe	Val	Ala	Arg	Asn	Arg	Ser	Ala	Cys	Thr	Tyr
				565						570					575	
	Leu	Ala	Ser	His	Leu	Arg	Gln	Leu	Gly	Cys	Ser	Pro	Asp	Leu	Arg	Gln
35				580						585					590	
	Arg	Leu	Arg	Tyr	Ala	Leu	Arg	Leu	Leu	Arg	Asp	Arg	Ser	Pro	Ala	
				595						600					605	

<210> 17

<211> 3492

5 <212> DNA

<213> Homo sapiens

<220> PTP-SL

<223> Description of Sequence: N/A

10

<400> 17

cagctaagac cgggagaggt ggaatttcac ttgaaattc ccttgccctg tgagggcccg 60
cgctgggcat gctcagtagc cgcggcgctg ctgctgggct gctgggctgg cgcggagtcc 120
accctgccgt ctccgccttg gcttctgggc gtccagaagg ccaggcattt gccgcctctg 180
15 agcgcttctg ttccccttac cgcgaacctc ctactgctct tcctctctcc ctctcttagg 240
gaggttgaag ctggtgctgg tttctgtcgg cgccacagac tgactgctct gcaaacccca 300
gccgaggacc tgaatcccgg agactagaag acccttggcg gtggctcttt ctaatagcac 360
tttacctgaa gtggggctcg ggtggagttt ctctccacc tctcaatgca aacactatgc 420
ggagagcagt ctgcttcctt gcgctgtgcc tgctccttaa tcttcacgct gcagggtgct 480
20 tttcaggaaa caatgatcat tttttggcaa ttaatcagaa gaagagtggg aagccgggtat 540
tcatttataa gcattcacia gacattgaga agagcctgga tatagcccca caaaaaatct 600
acagacatag ctaccattcc tcttccgaag ctcaagtaag caaacgccac cagattgtca 660
attcagcatt tcctagacct gcatatgacc cgtctctcaa tctgctggcc atggatggtc 720
aagatcttga agtggaaaat ctcccaatcc cagcagcaaa tgtaattgtg gtgacactgc 780
25 aaatggatgt aaacaagctg aacataacct tgcttcggat cttccgcca ggagtggctg 840
cagctttagg actcttacct cagcaagtgc acatcaatcg cctcattgga aagaagaaca 900
gtattgaact gtttgtgtct ccataaaacc gaaaaacagg aatttctgat gctctgccct 960
ctgaggaagt tcttcgttca cttaatatca atgttttgca tcaaagttaa tcccagtttg 1020
gaattacaga agtctctcct gagaaaaatg ttttacaagg gcagcatgaa gcggacaaaa 1080
30 tctggagcaa agaaggattt tatgctgttg tcatttttct cagcatcttt gttattatag 1140
taacgtgttt gatgattctt tacagattaa aagaaagatt tcagctttcc ttaagacaag 1200
acaaagagaa aaaccaggag atccacctat cgcccatcac attacagcca gcactgtccg 1260
aggcaaagac agtcacagc atggtccaac ctgagcaggc ccaaaggta ctgaatgttg 1320
tcgtggaccc tcaaggccga ggtgctcctg agatcagagc taccaccgct acctctgttt 1380
35 gcccttctcc tttcaaaatg aagcccatag gacttcaaga gagaagaggg tccaacgtat 1440
ctcttacatt ggacatgagt agcttgggga acattgaacc ctttgtgtct ataccaacac 1500
cacgggagaa ggtagcaatg gagtatctgc agtcagccag ccgaattctc acaaggtctc 1560
agctgaggga cgtcgtggca agttcacatt tactccaaag tgaattcatg gaaataccga 1620

```

tgaactttgt ggatcccaaa gaaattgata ttccgcgtca tggaactaaa aatcgctata 1680
agaccatttt accaaatccc ctcagcagag tgtgtttaag accaaaaaat gtaaccgatt 1740
cattgagcac ctacattaat gctaattata ttaggggcta cagtggcaag gagaaagcct 1800
tcattgccac gcagggcccc atgatcaaca ccgtggatga tttctggcag atggtttggc 1860
5 aggaagacag ccctgtgatt gttatgatca caaaactcaa agaaaaaat gagaaatgtg 1920
tgctatactg gccggaaaag agagggatat atggaaaagt tgaggttctg gttatcagtg 1980
taaataaatg tgataactac accattcgaa accttgtctt aaagcaagga agccacaccc 2040
aacatgtgaa gcattactgg tacacctcat ggcctgatca caagactcca gacagtgcc 2100
agccccctct acagctcatg ctggatgtag aagaagacag acttgcttcc cagggccgag 2160
10 ggcctgtggt tgtccactgc agtgcaggaa taggtagaac aggtgtgttt attgctacat 2220
ccattggctg tcaacagctg aaagaagaag gagttgtgga tgcactaagc attgtctgcc 2280
agcttcgtat ggatagaggt ggaatggtcc aaaccagtga gcagtatgaa tttgtgcacc 2340
atgctctgtg cctgtatgag agcagacttt cagcagagac tgtccagtga gtcattgaag 2400
acttgtcaga ccatcaatct cttggggtga ttaatcaaat taccacacca aggttcttag 2460
15 aaggagcttc ctgcaatgga aggaaggaga agctctgaag cccatgtatg gcatggattg 2520
tggaagactg ggcaacatat ttaagatttc cagctccttg tgtatatgaa tgcatttgta 2580
agcatcccc aaattattct gaaggttttt tgatgatgga ggtatgatag gttatcaca 2640
cagcctaagg cagattttgt tttgtctgta ctgactctat ctgccacaca gaatgtatgt 2700
atgtaatat cagtaataaa tgtcatcagg tgatgactgg atgagctgct gaagacattc 2760
20 gtattatgtg ttagatgctt taatgtttgc aaaatctgcc ttgtgaatgg actgtcagct 2820
gttaaactgt tcctgttttg aagtgtctat acctttctca gttaccagaa tcttgctgct 2880
aaagttgcaa gtgattgata atggattttt aacagagaag tctttgtttt tgaaaaacaa 2940
aaatcaaaaa cagtaactat tttatatgga aatgtgtctt gataatatta cctattaaat 3000
gtgtatttat agtccctcct atcaaacaat tacagagcac aatgattgtc attgggtata 3060
25 tatgtattta ctctctatta ttgggcataa aggtggett cgtccagaa ctctatccac 3120
tgtatttcca catcgtgagt cattttactt taaaaggga aaacaaattt gtagcaactc 3180
tgaagtatca agagttttaa ctacttgact ctcttttgct aagaagggat ttttgaatat 3240
gctatctacc tggaaatctc ctctcaacaa aaggtatatg ctttcaggaa tgatataatc 3300
tgtccattt tcgaggctcc ttataaggac atttccatgt atgtccttac atttctgaaa 3360
30 gctttcaatc ttcaagagcc aaaaaaatt aaaataacta ccctcagcaa acactagctg 3420
ttctgctcat atatgaattt ttaatgcagc aatgttgact ttgtttcata ctgccaataa 3480
actcttaata ct 3492

```

```

35 <210> 18
    <211> 657
    <212> PRT
    <213> Homo sapiens

```

<220>

<223> Description of Sequence: N/A

5 <400> 18

Met Arg Arg Ala Val Cys Phe Pro Ala Leu Cys Leu Leu Leu Asn Leu

1 5 10 15

His Ala Ala Gly Cys Phe Ser Gly Asn Asn Asp His Phe Leu Ala Ile

20 25 30

10 Asn Gln Lys Lys Ser Gly Lys Pro Val Phe Ile Tyr Lys His Ser Gln

35 40 45

Asp Ile Glu Lys Ser Leu Asp Ile Ala Pro Gln Lys Ile Tyr Arg His

50 55 60

Ser Tyr His Ser Ser Ser Glu Ala Gln Val Ser Lys Arg His Gln Ile

15 65 70 75 80

Val Asn Ser Ala Phe Pro Arg Pro Ala Tyr Asp Pro Ser Leu Asn Leu

85 90 95

Leu Ala Met Asp Gly Gln Asp Leu Glu Val Glu Asn Leu Pro Ile Pro

100 105 110

20 Ala Ala Asn Val Ile Val Val Thr Leu Gln Met Asp Val Asn Lys Leu

115 120 125

Asn Ile Thr Leu Leu Arg Ile Phe Arg Gln Gly Val Ala Ala Ala Leu

130 135 140

Gly Leu Leu Pro Gln Gln Val His Ile Asn Arg Leu Ile Gly Lys Lys

25 145 150 155 160

Asn Ser Ile Glu Leu Phe Val Ser Pro Ile Asn Arg Lys Thr Gly Ile

165 170 175

Ser Asp Ala Leu Pro Ser Glu Glu Val Leu Arg Ser Leu Asn Ile Asn

180 185 190

30 Val Leu His Gln Ser Leu Ser Gln Phe Gly Ile Thr Glu Val Ser Pro

195 200 205

Glu Lys Asn Val Leu Gln Gly Gln His Glu Ala Asp Lys Ile Trp Ser

210 215 220

Lys Glu Gly Phe Tyr Ala Val Val Ile Phe Leu Ser Ile Phe Val Ile

35 225 230 235 240

Ile Val Thr Cys Leu Met Ile Leu Tyr Arg Leu Lys Glu Arg Phe Gln

245 250 255

Leu Ser Leu Arg Gln Asp Lys Glu Lys Asn Gln Glu Ile His Leu Ser

	260		265		270
	Pro Ile Thr Leu Gln Pro Ala Leu Ser Glu Ala Lys Thr Val His Ser				
	275		280		285
	Met Val Gln Pro Glu Gln Ala Pro Lys Val Leu Asn Val Val Val Asp				
5	290		295		300
	Pro Gln Gly Arg Gly Ala Pro Glu Ile Arg Ala Thr Thr Ala Thr Ser				
	305		310		315
	Val Cys Pro Ser Pro Phe Lys Met Lys Pro Ile Gly Leu Gln Glu Arg				
		325		330	335
10	Arg Gly Ser Asn Val Ser Leu Thr Leu Asp Met Ser Ser Leu Gly Asn				
		340		345	350
	Ile Glu Pro Phe Val Ser Ile Pro Thr Pro Arg Glu Lys Val Ala Met				
		355		360	365
	Glu Tyr Leu Gln Ser Ala Ser Arg Ile Leu Thr Arg Ser Gln Leu Arg				
15	370		375		380
	Asp Val Val Ala Ser Ser His Leu Leu Gln Ser Glu Phe Met Glu Ile				
	385		390		395
	Pro Met Asn Phe Val Asp Pro Lys Glu Ile Asp Ile Pro Arg His Gly				
		405		410	415
20	Thr Lys Asn Arg Tyr Lys Thr Ile Leu Pro Asn Pro Leu Ser Arg Val				
		420		425	430
	Cys Leu Arg Pro Lys Asn Val Thr Asp Ser Leu Ser Thr Tyr Ile Asn				
		435		440	445
	Ala Asn Tyr Ile Arg Gly Tyr Ser Gly Lys Glu Lys Ala Phe Ile Ala				
25	450		455		460
	Thr Gln Gly Pro Met Ile Asn Thr Val Asp Asp Phe Trp Gln Met Val				
	465		470		475
	Trp Gln Glu Asp Ser Pro Val Ile Val Met Ile Thr Lys Leu Lys Glu				
		485		490	495
30	Lys Asn Glu Lys Cys Val Leu Tyr Trp Pro Glu Lys Arg Gly Ile Tyr				
		500		505	510
	Gly Lys Val Glu Val Leu Val Ile Ser Val Asn Glu Cys Asp Asn Tyr				
		515		520	525
	Thr Ile Arg Asn Leu Val Leu Lys Gln Gly Ser His Thr Gln His Val				
35	530		535		540
	Lys His Tyr Trp Tyr Thr Ser Trp Pro Asp His Lys Thr Pro Asp S r				
	545		550		555
	Ala Gln Pro Leu Leu Gln Leu Met Leu Asp Val Glu Glu Asp Arg Leu				

565 570 575
 Ala Ser Gln Gly Arg Gly Pro Val Val Val His Cys Ser Ala Gly Ile
 580 585 590
 Gly Arg Thr Gly Cys Phe Ile Ala Thr Ser Ile Gly Cys Gln Gln Leu
 5 595 600 605
 Lys Glu Glu Gly Val Val Asp Ala Leu Ser Ile Val Cys Gln Leu Arg
 610 615 620
 Met Asp Arg Gly Gly Met Val Gln Thr Ser Glu Gln Tyr Glu Phe Val
 625 630 635 640
 10 His His Ala Leu Cys Leu Tyr Glu Ser Arg Leu Ser Ala Glu Thr Val
 645 650 655
 Gln

15

<210> 19
 <211> 985
 <212> DNA
 <213> Homo sapiens

20

<220> HSP86
 <223> Description of Sequence: N/A

25

<400> 19
 ccggcccggt gtggctgtgc cggttggtcct gtgcgggtcac ttagccaaga tgccctgagga 60
 aaccagacc caagaccaac cgatggagga ggaggagggt gagacgttcg cctttcaggc 120
 agaaattgcc cagttgatgt cattgatcat caatactttc tactcgaaca aagagatctt 180
 tctgagagag ctcatctcaa attcatcaga tgcattggac aaaatccggt atgaaagctt 240
 gacagatccc agtaaattag actctgggaa agagctgcat attaacctta taccgaacaa 300
 30 acaagatcga actctcacta ttgtggatac tgggaattgga atgaccaagg ctgacttgat 360
 caataacctt ggtactatcg ccaagtctgg gaccaaagcg ttcattggaag ctttgcaggc 420
 tgggtgcagat atctctatga ttggccagtt cgggtgttgg ttttattctg cttattttggt 480
 tgctgagaaa gtaactgtga tcaccaaaca taacgatgat gagcagtacg cttggggagtc 540
 ctgagcaggg ggatcattca cagtgaggac agacacagggt gaacctatgg gtcgtggaac 600
 35 aaaagttatc ctacacctga aagaagacca aactgagtag ttggaggaac gaagaataaa 660
 ggagattgtg aagaaacatt ctgagtttat tggatatccc attactcttt ttgtggagaa 720
 ggaacgtgat aaagaagtaa gcgatgatga ggctgaagaa aaggaagaca aagaagaaga 780
 aaaagaaaaa gaagagaaag agtcggaaga caaacctgaa attgaagatg ttggttctga 840

tgaggaagaa gaaaagaagg atggtgacaa gaagaagaag aagaagatta aggaaaagta 900
catcgatcaa gaagagctca acaaaacaaa gcccatctgg accagaaatc ccgacgatat 960
tactaatgag gagtacggag aattc 985

5

<210> 20

<211> 312

<212> PRT

<213> Homo sapiens

10

<220>

<223> Description of Sequence: N/A

<400> 20

15 Met Pro Glu Glu Thr Gln Thr Gln Asp Gln Pro Met Glu Glu Glu Glu
1 5 10 15

Val Glu Thr Phe Ala Phe Gln Ala Glu Ile Ala Gln Leu Met Ser Leu
20 25 30

20

Ile Ile Asn Thr Phe Tyr Ser Asn Lys Glu Ile Phe Leu Arg Glu Leu
35 40 45

25

Ile Ser Asn Ser Ser Asp Ala Leu Asp Lys Ile Arg Tyr Glu Ser Leu
50 55 60

Thr Asp Pro Ser Lys Leu Asp Ser Gly Lys Glu Leu His Ile Asn Leu
65 70 75 80

30

Ile Pro Asn Lys Gln Asp Arg Thr Leu Thr Ile Val Asp Thr Gly Ile
85 90 95

Gly Met Thr Lys Ala Asp Leu Ile Asn Asn Leu Gly Thr Ile Ala Lys
100 105 110

35

Ser Gly Thr Lys Ala Phe Met Glu Ala Leu Gln Ala Gly Ala Asp Ile
115 120 125

Ser Met Ile Gly Gln Phe Gly Val Gly Ph Tyr Ser Ala Tyr Leu Val
 130 135 140

Ala Glu Lys Val Thr Val Ile Thr Lys His Asn Asp Asp Glu Gln Tyr
 5 145 150 155 160

Ala Trp Glu Ser Ser Ala Gly Gly Ser Phe Thr Val Arg Thr Asp Thr
 165 170 175

10 Gly Glu Pro Met Gly Arg Gly Thr Lys Val Ile Leu His Leu Lys Glu
 180 185 190

Asp Gln Thr Glu Tyr Leu Glu Glu Arg Arg Ile Lys Glu Ile Val Lys
 195 200 205

15 Lys His Ser Gln Phe Ile Gly Tyr Pro Ile Thr Leu Phe Val Glu Lys
 210 215 220

Glu Arg Asp Lys Glu Val Ser Asp Asp Glu Ala Glu Glu Lys Glu Asp
 20 225 230 235 240

Lys Glu Glu Glu Lys Glu Lys Glu Glu Lys Glu Ser Glu Asp Lys Pro
 245 250 255

25 Glu Ile Glu Asp Val Gly Ser Asp Glu Glu Glu Glu Lys Lys Asp Gly
 260 265 270

Asp Lys Lys Lys Lys Lys Lys Ile Lys Glu Lys Tyr Ile Asp Gln Glu
 275 280 285

30 Glu Leu Asn Lys Thr Lys Pro Ile Trp Thr Arg Asn Pro Asp Asp Ile
 290 295 300

Thr Asn Glu Glu Tyr Gly Glu Phe
 35 305 310